us-10-848-737-1.rge

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1706 bp RNA linear VRL 21-JUL-2003
SARS coronavirus Shanhgai LY orflab polyprotein and orfla
polyprotein genes, partial cds.
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HLKNGTCGLVELEKGYLPQLEQPYVPIKRSDALSTHYGHKVVELVABNDGIOYGRSGI
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WYTKHGSGALRELTRELNGGVYTRYDNNFCGPDGYPLDCIKDFLABAGKSMCTLSED
LDYIESKRGVYCCRDHEHEINGWYTRYDNNFCGPDGYPLDCIKDFLARAKKFDTFKGECPKFVFP
LNSKYKVIQEYVEKKKTEGFWGRIRSYYPVASPQECNNWHLSTJMKCNHCDBYSWQTC
DFLKATCEHCGTENLVIEGFTTCGYLPTNAVVKMPFDACQDPEIGPENSVADYNHSN
IETRLRKGGFRFCFGGCVFAYVGCYNKRAYWPRASADIGSGHTGTTGDNVETINEDL
LELLSRERNININGDFHLNEEVAIILASFSASTSAFIDTIKSLDYKKFFTINEDC
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SARS coronavirus Shanhgai LY
SARS coronavirus Shanhgai LY
Viruses; seRNA positive—strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
1 (bases 1 to 1706)
1 (bases 1 to 1706)
1 (bases 1 to 1706)
Vian, Z., Zhang, X., Hu, Y., Lan, S., Wang, H., Zhou, Z. and Wen, Y.
Direct Submission
Submitted (12-JUN-2003) Molecular Virology, Shanghai Medical
College of Fudan University, 138 Yi Xue Yuan Road, Shanghai 200032,
P.R. China
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/note="translated by ribosome slippage event"
/codon_start=1
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                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
1. 1706
| organism="SARS coronavirus Sh|
|mol type="genomic RNA" |
|db_xref="taxon:235173" |
|country="China: Shanghai" |
|186. .>1706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /product="orfla polyprotein"
/protein_id="AAP82966.1"
/db_xref="GI:32454344"
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                                                     June 23, 2005, 23:08:06; Search time 763.417 Seconds (without alignments) 1396.372 Million cell updates/sec
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      GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Wang, Z.G., Li,L.J., Luo,Y., Zhang,J.Y., Wang,M.Y., Cheng,S.Y., Zhang,Y.J., Wang,X.M., Lu,Y.Y., Wu,N.P., Mei,L.L. and Wang,Z.X. Molecular biological analysis of genotyping and phylogeny of severe accute respiratory syndrome associated coronavirus Chin. Med. J. 117 (1), 42-48 (2004)
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TGVLVPHVDGEPIAYRNVLLKKNGRKGAGGHSVGIDLKSYDGJBELFTPDIBDYBQN
WNTKHGSGALRELTEBLNGGAVTRVVDNHGCGPDGYPLDCI KOPLARAGKSMCTLSBG
LDYIESKRGYVCCRDHEHBIAWFTBRSDKSYEHQTPPEIKSAKKFDTFKGECPKFVPP
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DETAATCHGTGTENLVIEGPTTGGTLPTNAVVRAPCPACOPDEIGPEHSVADYTHNEN
IETRLKKGRTRCFGGCVPAYVGCYNKRAYWPRASADIGSGHTGITGDNVTLNEDL
LEILESRERVNINIVGDFHLNEEVAIILASFSASTSAFIDTIKSLDYKSFKTIVESCGN
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                                           Viruses; ssRNA positive-strand viruses, no DNA stage; Nidovirales; Coronaviridae; Coronavirus.

1 (bases 1 to 2633)

Li,L., Wang,Z., Lu,Y., Bao,Q., Chen,S., Wu,N., Cheng,S., Weng,J., Zhang,Y., Yan,J., Mei,L., Wang,X., Zhu,H., Yu,Y., Zhang,M., Li,M., Sao,J., Mei,L., Wang,X., Wang,S. and Hu,S. Severe acute respiratory syndrome-associated coronavirus genotype and its characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /isolation source="cultured virus in Vero cell inoculated with throat swab from patient with severe acute respiratory syndrome (SARS)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (13-NOV-2003) Department of Microbiology, Zhejiang CDC, Submitted (13-NOV-2003) Department of Microbiology, Zhejiang CDC, 17 Laozhedazhi Rd., Hangzhou, Zhejiang 310009, China Equence update by submitter.

6 (bases 1 to 2633)

8 Mang, Z., Cheng, S. and Zhang, Y.

10 Irect Submission

Submitted (17-Daozhedazhi Rd., Hangzhou, Zhejiang 310009, China Sequence update by submitter.

Sequence update by submitter.

17 Laozhedazhi Rd., Hangzhou, Zhejiang 310009, China Sequence update by submitter.

18 Sequence version replaced gi:38304880.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wang, Z., Cheng, S. and Zhang, Y.
Direct Submission

17 Laozhedazhi Rd., Hangzhou, Zhejiang 310009, China
4 (Dases I to 26333)
Wang, Z., Cheng, S. and Zhang, Y.
Direct Submission
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    1. .248
    51. .56
    /note="transcription regulatory">/note="transcription regulatory

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/country="China: Hangzhou"
/note="isolated April 21, 2003"
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/note="ribosomal slippage"
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/db_xref="G1:38324306"
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ng, Z., Cheng, S. and Zhang, Y.
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SARS coronavirus ZJ01
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LDYI BSKRGYYCRDHEHELIAWFTERSDKSCEHQTPFEIKSAKKFDTFKGECPKFVTP
LNSKVKVI OPRVEKKKTGFRØRS IRSYYPVAS POECHWHLSTLMKCHHODEVSWQTC
DFLKATCEHCGTENLVIEGPTTCGYLPTNAVVKMPCPACQDPEIGPEHVADINHSKN
                                                                                                                                                                                                                                                            IETRLRKGGRTRCFGGCVFAYVGCYNKRAYWVPRASADIGSGHTGITGDNVETLNEDL
LEILSRERVNINIVGDFHLNEEVAIILASFSASTSAFIDTIKSLDYKSFKTIVESCG"
                                                                                                 HLKNGTCGLVELEKGVLPQLEQPYVFIKRSDALSTNHGHKVVELVAEMDGIQYGRSGI
TLGVLVPHVGETPIAYRNVLLRKNGNKGAGGHSYGIDLKSYDLGDELGTDPIEDYEQN
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                                                                            /translation="MESLVLGVNEKTHVQLSLPVLQVRDVLVRGFGDSVERALSBARE
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SARS coronavirus Sin0409
Viruses; ssRNA positive-strand viruses, no DNA stage; Nidovirales;
Viruses; csRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus
Lim.P.L., Kurup.A., Gopalakrishna,G., Chan,K.P., Wong,C.W.,
Ng,L.C., Se-Thoe,S.Y., Oon,L., Bai,X., Stanton,L.W., Ruan,Y.,
Miller,L.D., Vega,V.B., James,L., Ooi,P.L., Kai,C.S., Olsen,S.J.,
Ang,B. and Leo,Y.S.
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Wei,C.L., Lee,C., Lin,S., Thoreau,H., Vega,V.B., Stanton,L.W. and
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Submitted (28-JAN-2004) Genome Institute of Singapore,
Street, 02-01, Genome, Singapore 138672, Singapore
Location/Qualifiers
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SARS coronavirus Sin0409, partial sequence.
/product="orflab polyprotein"
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SARS coronavirus ZJ01, partial genome.
AY286320 AY286320.4 GI:39980888
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/country="Singapore"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    697 GTGAACTCACTCGTGAGCTC 716
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231 GTGAACTCACTCGTGAGCTC 250
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Best Local Similarity 75.0%;
Matches 15; Conservative
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Best Local Similarity 75.0
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AY286320
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sequence leader; TRS"

EAARCMRSLKAPAVVSVSSPDAVTTYNGYLTSSSKTSEEHFVETVSLAGSYRDWSYSG QRTELGVEFLKRGDK LYYHTLESPVEFHLIOGEVLSLDKLKSLLSLREVKTIKVFTTVD YTTLHTQLVDWSMTYGQQFGPTVLDGADVTK IKPHVNHEGKTFFVLPSDDTLRSBAFE YYHTLDESFLGRYWARLAHTKKWKFPQVGGITS IKMADNNCYLSSVLLALQDLEVKRN APALQEAYYRARAGDAANFCALILAYSNKTVGELGDVRETWTHLLQHANLESBAKRVLN AKKCLHVVGPNLNAGEDI QLLKAAYENFNSQDI LLAPLLSAGI FGAKPLOSLOVCVOT VRTQVY I AVNDKALYEQVVMDYLDNLKPRVEA PKQEEPPNTEDSKTEEKS VVQKPVDV KPKI KACI DEVTTTILEETKFLTNKLLLFAD I NGKLYHDSQNMLRGEDMSFLEKDAPYM GAKLLHKPIVWHINQATTKTTFKPNTWCLRCLWSTKPVDTSNSFEVLAVEDTQCMDNL ACESQQPTSEEVVENPTIQKEVIECDVKTTEVVGNVILKPSDEGVKVTQELGHEDLMA AYVENTSITIKKPNELSLALGLKTIATHGIAAINSVPWSKILAYVKPFLGQAAITTSN CAKRLAQRVFUNYMPYVFTLLFQLCTFTKSTNSRIRASLPTTIAKNSVKSVAKLCLDA GINYVKSPKFSKLFTIAMMLLLLSICLGSLICVTAAFGVLLSNFGAPSYCNGVRELYL NSSNVTTMDFCEGSFPCSICLSGLDSLDSYPALETIQVTISSYKLDLTILGLAAEWVL ASFYYIWKSYVHIMDGCTSSTCMMCYKRNRATRVBCTTIVNGMKRSFYVYANGGRGFC KTHNWNCLNCDTFCTGSTFISDEVARDLSLQFKRPINPTDQSSYIVDSVAVKNGALHL YFDKAGQKTYERHPLSHFVNLDNLRANNTKGSLPINVIVFDGKSKCDESASKSASVYY HTLSIHDGYTNEIIGYKAIQDGVTRDIISTDDCFANKHAGFDAWFSQRGGSYKNDKSC PVVAAIITREIGFIVPGLPGTVLRAINGPFLHFLPRVFSAVGNICYTPSKLIEYSDFA TSACVLAAECTIFKDAMGKPVPYCYDTNLLEGSISYSELRPDTRYVLMDGSIIQFPNT Y CALS PGLLATNINV FRLKGGAP I KGVT FGED TVWEV OG Y KNVR I TFELDER VDKVLNE KCSVYTVESGTEVTEFACVVAEAVVKTLQPVSDLLTNMG1DLDEWSVATFYLFDDAGE ENFSSRMYCSFYPPDEEEEDDAECEEEEIDETCEHEYGTEDDYQGLPLEFGASAETVR VEEEEEEDWLDDTTEQSEIEPEPPPEPVNQFTGYLKLTDNVAIKCVDIVKEAQSA NPMVIVNAANIHLKHGGGVAGALNKATNGAMQKESDDYIKLNGPLTVGGSCLLSGHNL QRKYKGIKIQEGIVDYGVRFFPYTSKEPVASIITKLNSLNEPLVTMPIGYVTHGFNLE /VCKHCGQKTTTLTGVEAVMYMGTLSYDNLKTGVSIPCVCGRDATQYLVQQESSFVMM SAPPAEYKLQQGTFLCANEYTGNYQCGHYTH1TAKETLYRIDGAHLTKMSEYKGPVTD VFYKETSYTTTIKPVSYKLDGVTYTE1EPKLDGYYKKDNAYYTEQP1DLVPTQPLPNA SFDNFKLTCSNTKFADDLNQMTGFTKPASRELSVTFFPDLNGDVVAIDYRHYSASFKK AYMLFTKFFYLLGLSAIMQVFFGYFASHFISNSWLMWFI1SIVQMAPVSAMVRMYIFF SQLMCQP1LLLDQALVSDVGDSTEVSVKMFDAYVDTFSATFSVPMEKLKALVATAHSE LAKGVALDGVLSTFVSAARQGVVDTDVDTKDV1ECLKLSHHSDLEVTGDSCNNFMLTY NKVENMTPRDLGACIDCNARHINAQVAKSHNVSLIWNVKDYMSLSEQLRKQIRSAAKK NNIPPRLTCATTRQVVNVITTKISLKGGKIVSTCFKLMLKATLLCVLAALVCYIVMPV TLEGSVRVVTTFDAEYCRHGTCERSEVGICLSTSGRWVLNNEHYRALSGVFCGVDAMN LIANIFTPLVQPVGALDVSASVVAGGIIAILVTCAAYYFMKFRRVFGEYNHVVAANAL LFLMSFTILCLVPAYSFLPGVYSVFYLYLTFYFTNDVSFLAHLQWFAMFSPIVPFWIT AIYVFCISLKHCHWFFNNYLRKRVMFNGVTFSTFEEAALCTFLLNKEMYLKLRSETLL TSAVLQSGFRKMAFPSGKVEGCMVQVTCGTTTLNGLWLDDTVYCPRHV1CTAEDMLNP TFSVLACYNGSPSGVYQCAMRPNHT1KGSFLNGSCGSVGFN1DYDCVSFCYMHHMELP TGVHAGTDLEGKFYGPFVDRQTAQAAGTDTTTTLAVLAMLYAAVINGDRWFLNRFTTT LNDFNLVAMKYNYEPLTQDHVDILGPLSAQTGIAVLDMCAALKELLQNGMNGRTILGS FKLNI KLLGI GGKPCI KVATVQSKMSDVKCTSVVLLSVLQQLRVBSŠSKLMAQCVQLH NDI LLAKDTTEAFEKMVSLLSVLLSMQGAVDI NRLCEEMLDNRATLQAI ASEFSSLPS YAAYATAQEAYEQAVANGDSEVVLKKLKKSLMVAKSEFDRDAAMQRKLEKMADQAMTQ MYKQARSEDKRAKVTSAMQTMLFTMLRKLDNDALNNI INNARDGCVPLNI I PLTTAAK **ISTDVVYRAFDIYNEKVAGFAKFLKTNCCRFQEKDEEGNLLDSYFVVKRHTMSNYOHE** GIVGVLTLDNODLNGNWYDFGDFEQVAPGCGVPIVDSYYSLLMPILTLTRALAAESHM DADLAKPLIKWDLLKYDFTBERLCLFDRYFKYWDQTYHPNCINCLDDRCILHCANFNV LFSTVFPPTSFGPLVRKIFVDGVPFVVSTGYHFRELGVVHNQDVNLHSSRLSFKELLV NKALEMCIDQVTIAGAKLRSLNLGEVFIAQSKGLYRQCIRGKEQLQLLMPLKAPKEVT FLEGDSHDTVLTSEEVVLKNGELEALETPVDSFTNGAIVGTPVCVNGLMLLEIKDKEQ /GDVITSGDITCVVIPSKKAGGTTEMLSRALKKVPVDEYITTYPGQGCAGYTLEEAKT NYEDLLIRKSNHSFLVQAGNVQLRVIGHSMQNCLLRLKVDTSNPKTPKYKFVRIQPGQ TILEDEFTPFDVVRQCSGVTFQGKFKKIVKGTHHWMLLTFLTSLLILVOSTOWSLFFF **VYENAFLPFTLGIMAIAACAMLLVKHKHAFLCLFLLPSLATVAYFNMVYMPASWVMRI** MTWLELADTSLSGYRLKDCVMYASALVLLILMTARTVYDDAARRVWTLMNVITLVYKV MLVYCFLGYCCCCYFGLFCLLNRYFRLTLGVYDYLVSTQEFRYMNSQGLLPPKSSIDA LMVVVPDYGTYKNTCDGNTFTYASALWEIQQVVDADSKHVQLSEINMDNSPNLAWPLI VTALRANSAVKLQNNELSPVALRQMSCAAGTTQTACTDDNALAYYNNSKGGRFVLALL SDHQDLKWARFPKSDGTGTIYTELEPPCRFVTDTPKGPKVKYLYFIKGLNNLNRGMVL GSLAATVRLQAGNATEVPAHSTVLSFCAFAVDPAKAYKDYLASGGQPITNCVKMLCTH TGTGQAITVTPEANMDQESFGGASCCLYCRCHIDHPNPKGFCDLKGKYVQIPTTCAND PVGFTLRNTVCTVCGMWKGYGCSCDOLREPLMOSADASTFLNRVCGVSAARLTPCGTG ALKKCKSAFYVLPSEAPNAKEEILGTVSWNLREMLAHAEETRKLMPICMDVRAIMATI PLTQYNRYLALYNKYKYFSGALDTTSYREAACCHLAKALNDFSNSGADVLYQPPQTSI YYGNALDQAISMWALVISVTSNYSGVVTTIMFLARAIVFVCVEYYPLLFITGNTLQCI ETIYNLVKDCPAVAVHDFFKFRVDGDMVPHISRQRLTKYTMADLVYALRHFDEGNCDT

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IVVPDEISMATNYDLSVVNARLRAKHYYYIGDPAQLPAPRILITKGTLEPEYFNSVCR
LMKTIGPDMFLGTCRCCPAEIVDTVSALVYDNKLKAHKDKSAQCFKMFYKGVITHDVS
SAINRPQIGVVREFLTRNPAMRKAVFISPYNSONASKILGLFTQTVDSSQSSEYDY
STITTGTTGTHYNVAITRAKFIGILCIMSDRDLYDKLGFTSLEIPRRNYATLQ
ABNYTGLFKDCSKITTGLHPTQAPTHLSVDIKKTEGLCVDIFGIPKDMTYRRISSM
GFKMNYQVNGYPNMFITREEAIRHVRAMIGFDVEGCHATRDAVGTNLPLDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LVAVPTGYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPMNVVRIKIVQMLSDTL
KGLSDRVVFVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACMNHSVG
FDYVYNPFMIDVQQMGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRV
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HLKNGTCGLVblekGvlpQleQpYvFlkRSDalsTnHGHKvVelvaemDGIQYGRSGI
TLGVLVPHVGETPIAYRNVLLRKNGNKGAGGHSYGIDLKSYDLGDELGTDPIEDYEQN
                                                                                                               KNFKAVLYYQNNVFMSEAKCWTETDLTKGPHEFCSQHTWLVKQGDDYVYLPYPDPSKI
LGAGCFVDDIVKTDGTLMIERFVSLAIDAYPLTKHPNQEYADVFHLYLQYIRKLHDEL
TGHMLDMYSVMLTNDNTSRYWEPEFYEAMYTPHTVLQAVGACVLCNSQTSLRCGACIR
                                                                                                                                                                                                                                            SFPLCANGQVFGLYKNTCVGSDNVTDFNAIATCDWTNAGDYILANTCTERLKLFAAET
LKATEETFKLSYGIATVREVLSDRELHLSWEVGKPRPPLNRNYVFTGYRVTKNSKVQI
                                                                                                                                                                                                                                                                                                                                  PTLNI SDEFSSNVANYQKVGMQKYSTLQGPPGTGKSHFA I GLALYYPSAR I VYTACSH
AAVDALCEKALKYLPIDKCSR I I PARARVECFDKFKVNSTLEQYVFCTVNALPETTAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KFYDAQPCSDKAYKI EELFY SYATHHDKFTDGVCLFWNCNVDRY PANA I VCRFDTRVL
SNINLPGCDGGSLYVNKHA FHTPAFDKSAFTNLKOLPFYYSDSPCESHGKOVVSDID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FTRLQSLENVAYNVOKGHFDGHAGEAPVSIINNAVYTKVDGIDVEIFENKTTLPVNV
AFELWAKRNIKPVPEIKILNNLGVDIAANTVIWDYKREAPAHVSTIGVCTWTDIAKKP
TESACSSLTVLFDGRVEGQVDLFRNARNGVLITEGSVKGLTPSKGPAQASVNGVTLIG
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GSSKCVCSVIDLLLDDFVEIIKSQDLSVISKVVKVTIDYAEISFMLWCKDGHVEFFYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KLQASQAWQPGVAMPNLYKMQRMLLEKCDLQNYGENAVI PKGIMMNVAKYTQLCQYLN
TLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQMLPTGTLLVDSDLNDFVSDADSTLIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FWRNTNPIQLSSYSLFDMSKFPLKLRGTAVMSLKENQINDMIYSLLEKGRLIIRENNR
                            YSDVETPHLMGWDYPKCDRAMPNMLRIMASLVLARKHNTCCNLSHRFYRLANECAQVL
                                                      SEMVMCGGSLYVKPGGTSSGDATTAYANSVFNICQAVTANVNALLSTDGNKIADKÝVR
NLQHRLYECLYRNRDVDHEFVDEFYAYLRKHFSMMILSDDAVVCYNSNYAAQGLVASI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DWSVEYPIIGDELRVNSACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQAEVEW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YVPLKSATCITRCNLGGAVCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNT
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ISAKNRARTVAGVSICSTMTNRQFHQKLLKSIAATRGATVVIGTSKFYGGWHNMLKTV
                                                                                                                                                                                                                                                                                                         GEYTFEKGDYGDAVVYRGTTTYKLNVGDYFVLTSHTVMPLSAPTLVPQEHYVRITGLY
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                                                                                                                                                                                                            RPFLCCKCCYDHVISTSHKLVLSVNPYVCNAPGCDVTDVTQLYLGGMSYYCKSHKPPI
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SARS coronavirus ShanghaiQXC2
Viruses; BRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
I (bases 1 to 29013)
Yuan, Z., Zhang, X., Hu, Y., Lan, S., Zhou, Z., Wang, H. and Wen, Y.
Analysis of SARS coronavirus genome in Shanghai isolates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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'protein id="AAR16181.1"
'db_xref="GI:38304882"
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RPI FEWI EAKLSAGVEFLKDAWEI LKPLITGVFDI VKGQI QVASDNI KDCVKCFI DVV
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FLEGDSHDTVLTSEEVVLKNGELEALETPVDS FTNGAI VGTPVCVNGLMLLEI KDKEQ
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VVCKHCGQKTTTLTGVEAVMYMGTLSYDNLKTGVSIPCVCGRDATQYLVQQESSFVMM
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YKVTKGKPVKGAMNI GQQRSVLTPLCGFPSQAAGVI RSI FARTLDAANHSI PDLQRAA
VTI LDGI SEQSLRLVDAMVYTSDLLTNSVI IMAYVTGGLVQQTSQMLSNLLGTTVEKL
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AKKCLHVVGPNLNAGEDI QLLKAAYENFNSQDI LLAPLLSAGI FGAKPLGSLQVCVQT
VRTQVY I AVNDKALYEQVVMDYLDNLKPRVEAPKQEEPPNTEDSKTEEKSVVQKPVDV
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LLFQLCTFTKSTNSRIRASLPTTIAKNSVKSVAKLCLDAGINYVKSPKFSKLFTIAMW
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RHINAQVAKSHNVSLI WNVKDYMSLSEQLRKQI RSAAKKNNI PFRLTCATTRQVVNVI
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WNTKHGSGALRELTRELNGGAVTRYVDNNFCGPDGYPLDCIKDFLARAGKSMCTLSEQ
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DFLKATCEHCGTENLVI EGPTTCGYL PTNAVVKMPCPACQDPE I GPEHSVADYHNHSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    I ETRLIRKGGRTRCFGGCVFAYVGCYNKRAYWVPRASAD I GSGHTGI TGDNVETLNEDL
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NLDNLRANNTKGSLPINVI VFDGKSKCDESASKSASVYYSQLMCQPILLIDQALVSDV
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                                                     2 (bases 1 to 29013)
Yuan, Z., Zhang, X., Hu, Y., Lan, S., Zhou, Z., Wang, H. and Wen, Y.
Direct Submission
Submitted (11-NOV-2003) Key Lab of Medical Molecular Virology,
Shanghai Medical College, Fudan University, 138 Yi Xue Yuan Road,
Shanghai 200032, P.R. China
                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="SARS coronavirus Shanghai QXC2"
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CKYTQSTALQELLIQQMIQFMMSRRRLLACLCKHKKVSTNLCTHSFRKKQVR"
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FCTQLNRALSGIAAEQDRNTREVFAQVKQMYKTPTLKYFGGFNFSQILPDPLKPTKRS
FIEDLLFNKVTLADAGFMKQYGECLGDINARDLICAQKFNGLTVLPPLLTDDMIAAYT
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From independent foci of epidemic outbreak to large genomic
alteration in late phase viruses: evolution of the SARS-coronavirus
                   FEYISDAFSLDVSEKSGNFKHLREFVFKNKDGFLYVYKGYQDIDVVRDLPSGFNTLKP
I FKLPLGINITNFRAILTAFSPAQDIMGTSAAAXFVGYLKPTTFMLKYDENGTITDAV
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Coronaviridae; Coronavirus.
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AY394999
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product="unknown"
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Pred. No. 6.5;
5; Mismatches 0; Indels 0;
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Submitted (19-SEP-2003) Guangdong, China
Location/Qualifiers
1. . 29350
/organism="SARS coronavirus LC2"

    .29350
/organism="SARS coronavirus LC3"

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SARS coronavirus LC4, complete genome.
AY395001
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                                                                                 /mol_type="genomic_RNA"
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/db_xref="taxon:249084"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="genomic RNA"
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/db_xref="taxon:249085"
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761 GTGACTCACTCGTGAGCTC 780
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761 GTGAACTCACTCGTGAGCTC 780
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llarity 75.0%;
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SARS coronavirus LC3
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nes 15; Conserv
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SARS coronavirus HSZ-Bb
Vituese; sRRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronavirides; Coronavirus.
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SARS coronavirus Taiwan TC1, complete genome.
AY338174 GI:32493129
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/isolate="HSZ-Bb"
/db_xref="taxon:249070"
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/organism="SARS coronavirus GZ-A"
/nol type="genomic RNA"
/isolate="GZ-A"
/db_xref="taxon:249062"
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AY394985
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526 GTGAACTCACTCGTGAGCTC 545
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725 GTGAACTCACTCGTGAGCTC 744
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Best Local Similarity 75.0
Matches 15; Conservative
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Matches 15; Conservative
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AY338174
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SARS coronavirus LC5
SARS coronavirus LC5
Viruses; saRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
1 (bases 1 to 293S0)
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SARS coronavirus G2-A
Viruses; sBRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
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                                                                                                                                /organism="SARS coronavirus LC4"
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/db_xref="taxon:249086"
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/mol_type="genomic RNA"
/isolate="LC5"
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SARS coronavirus GZ-A, partial genome.
AY394977
AY394977.1 GI:37624320
                                                                                                                                                                                                                                                                                                                                                                                                                    RNA
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AY395002
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Best Local Similarity 75.0%; Pred. No. 6
Matches 15; Conservative 5; Mismatch
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IETRLRKGGRTRCFGGCVFAYVGCYNKRAYWVPRASADIGSGHTGITGDNVETLNEDL
LEILSRERVNINIVGDFHLNEEVAIILASFSASTSAFIDTIKSLDYKSFKTIVESCGN
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VTILDGISEQSLRLVDAMVYTSDLLTNSVIIMAYVTGGLVQQTSQWLSNLLGTTVEKL
RPIFEWIEAKLSAGVEFLKDAMEILKPLITGVPDIVKGQIQVASDNIKDCVKCPIDVV
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VEEEEEEDWLDDTTEGSEIEPEPPTPEEPVNOFTGYLKLTDNVAIKCVDIVKEAQSA
NPMVIVNAANIHLKHGGGVAGALNKATNGAMQKESDDYIKLNGPLTVGGSCLLSGHNL
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                                SARS coronavirus Taiwan TC1
SARS coronavirus Taiwan TC1
Viruses; ssRNA positive-strand viruses, no DNA stage; Nidovirales;
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Submitted (08-JUL-2003) Department of Molecular Medicine, China Medical University Hospital, 2, Yuh Der Road, Taichung, Taichung 404, Taiwan to 29573)
3 (bases I to 29573)
Chang,J.-G.C., Lin,T.-H., Chen,C.-M., Lin,C.-S., Chan,W.-L. and
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404, Taiwan
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produc=**nonstructural polyprotein pplab**
/protein_id=**AAQ01594.1**
/db_xref=**GI:33285967**
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organism="SARS coronavirus Taiwan TC1"
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note="clinical specimen"
oin(196. .13329,13329. .21416)
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isolate="TC1"
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Direct Submission
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LIANI FTPLVQPVGALDVSASVVAGGI IAILVTCAAYYFMKFRRVFGEYNHVVAANAL LFLMSFTILCLVPRYSFLPGYYSVFTLYLTFYFTNDVSFLAHLØMFAMFSPIVPFWIT YTVFCI SIKHCHWFFNYTRRRVMFNGVTFSTFEBAALCTFLLMKEMYLKLRSFTLL PLTQYNRYLALYNKYKFSGALDTTSYRBAACCHLAKALNDFSNSGADVLYQPOPQTSI TSAVLQSGFRKMAFPSGKVEGCMVQVTCGTTTLNGLWLDDTVYCPRHVI CTAEDMLNP CAKRLAQRVFNNYMPYVFTLLFQLCTFTKSTNSRIRASLPTTIAKNSVKSVAKLCLDA GINYVKSPKFSKLFTIAMMLLLLSICLGSLICVTAAFGVLLSNPGAPSYCNGVRELYL NSSNVTTMDFCEGSFPCSICLSGLDSLDSYPALETIQVTISSYKLDLTILGLAAEWVL SQLMCQPILLLDQALVSDVGDSTEVSVKMFDAYVDTFSATFSVPMEKLKALVATAHSE LKKYALDGYLSTFVSAARGKYVDTVDTKDVI ECLKLSHHSDLEVTGSCNNFKLTY INVENMTPRDLGACIDCNARHINAQVAKSHVYSLIWYKDYMSLSEQLRKQIRSAAKK NNI PPRLCATTRQVVNVITTKISLKGGKIVSTCKKAMKATLLCVLAALVCYIVMPV TGVHAGTDLEGKFYGPFVDRQTAQAAGTDTT1TLNVLAMLYAAVINGDRWFLNRFTTT LNDFNLVAMKYNYEPLTQDHVDILGPLSAQTGIAVLDMCAALKELLQNGMNGRTILGS TILEDEFTPFDVVRQCSGVTFQGKFKKIVKGTHHWMLLFFLTSLLILVQSTQWSLFFF VYENAFLPFTLGIMAIAACAMLLVKHKHAFLCLFLLPSLATVAYFNMVYMPASWVMRI MTWLELADTSLSGYRLKDCVMYASALVLLILMTARTVYDDAARRWTLMNVITLVYKV MLVYCFLGYCCCCYFGLFCLLMRYFRLTLGVYDYLVSTQEFRYMNSQGLLPPKSSIDA FKLNIKLLGIGGKPCIKVATVQSKMSDVKCTSVVLLSVLQQLRVESSSKLMAQCVQLH NDILLAKDTTEAFEKMVSLLSVLLSMQGAVDINRLCEEMLDNRATLQAIASEFSSLPS GIVGVLTLDNQDLNGNWYDFGDFVQVAPGCGVPIVDSYYSLLMPILTLTRALAAESHM DADLAKPLIKWDLLKYDFTEERLCLFDRYFKYWDQTYHPNCINCLDDRCILHCANFNV LFSTVFPPTSFGPLVRKIFVDGVPFVVSTGYHFRELGVVHNQDVNLHSSRLSFKELLV YSDVETPHLMGWDYPKCDRAMPUMLRIMASLVLARKHNTCCNLSHRFYRLANECAQVL SEMVMCGGSLYVKPGGTSSGDATTAYANSVFNICQAVTANVNALLSTDGNKIADKYVR NLQHRLYECLYRNRDVDHEFVDEFYAYLRKHFSMMILSDDAVVCYNSNYAAQGLVASI AAVDALCEKALKYLPIDKCSRIIPARARVECFDKFKVNSTLEQYVFCTVNALPETTAD IVVFDEISMATNYDLSVVNARLRAKHYVYIGDPAQLPAPRTLLTKGTLEPEYFNSVCR LMXTIGPDMFLGTCRRCPAEIVDTVSALVYDNKLKAHKDKSAQCFKMFYKGVITHDVS AYMLPTKFFYLLGLSAIMQVFFGYFASHFISNSWLMWFIISIVOMAPVSAMVRMYIFF SEYYIWKSYYHIMDGCTSSTCMCYKRNRATHYEVCTTIVMGMRSFYYANGGRGFC KTHWNCINCINCOFFGSTFISDEVARDI.SLOFKRPINFTDQSSYIVDSVAVKNGALHL YFDKAGQKTYERHPLSHFVNLDNLRANNTKGSLPINVIVFDGKSKCDESASKSASVYY TSACVLAAECTIFKDAMGKPVPYCYDTNLLEGSISYSELRPDTRYVLMDGSIIQFPNT YLEGSVRVVTTFDAEYCRHGTCERSEVGICLSTSGRWVLNNEHYRALSGVFCGVDAMN NYEDLLIRKSNHSFLVQAGNVQLRVIGHSMQNCLLRLKVDTSNPKTPKYKFVRIQPGQ TFSVLACYNGSPSGVYQCAMRPNHTIKGSFLNGSCGSVGFNIDYDCVSFCYMHHMELP LMVVVPDYGTYKNTCDGNTFTYASALWEIQQVVDADSKIVQLSEINMDNSPNLAWPLI VTALRANSAVKLQNNELSPVALRQMSCAAGTTQTACTDDNALAYYNNSKGGRFVLALL GSLĀATVRLQAGNATEVPANSTVLSFCAFAVDPAKAYKDYLASGGQPITNCVKMLCTH TGTGQAITVTPEANMDQESFGGASCCLYCRCHIDHPNPKGFCDLKGKYVQIPTTCAND PVGFTLRNTVCTVCGMMKGYGCSCDQLREPLMQSADASTFLNRVCGVSAARLTPCGTG TSTDVVYRAFDIYNEKVAGFAKFLKTNCCRFQEKDEGGNLLDSYFVVKRHTMSNYQHE ETIYNLVKDCPAVAVHDFFKFRVDGDMVPHISRQRLTKYTMADLVYALRHFDEGNCDT LKEILVTYNCCDDDYFNKKDWYDFVENPDILRVYANLGERVRQSLLKTVQFCDAMRDA GSSVELKHFFFAQDGNAAISDYDYYRYNLPTMCDÎRQILFVVEVVDKYFDCYDGGCIN ANQVIVNNLDKSAGFPFNKWGKARLYYDSMSYEDQDALFAYTKRNVIPTITQMNLKYA PVVAAIITREIGFIVPGLPGTVLRAINGDFLHFLPRVFSAVGNICYTPSKLIEYSDFA YYGNALDQAISMWALVISVTSNYSGVVTTIMFLARAIVFVCVEYYPLLFITGNTLQCI YAAYATAQEAYEQAVANGDSEVVLKKUKKSLNVAKSEFDRDAAMQRKLEKMADQAMTQ MYKQARSEDKRAKVTSAMQTMLFTMLRKLDNDALNNI INNARDGCVPLNI IPLTTAAK SDHQDLKWARFPKSDGTGTIYTELEPPCRFVTDTPKGPKVKYLYFIKGLNNLNRGMVL YAADPAMHAASGNLLLDKRTTCFSVAALTNNVAFQTVKPGNFNKDFYDFAVSKGFFKE KNFKAVLYYQNNVFMSEAKCWTETDLTKGPHEFCSQHTMLVKQGDDYVYLPYPDPSR.I LGAGCFVDDIVKTDGTLMIERFVSLAIDAYPLTKHPNQEYADVFHLYLQYIRKLHDEL TGHMLDMYSVMLTNDNTSRYWEPEFYEAMYTPHTVLQAVGACVLCNSQTSLRCGACIR SFPLCANGOVFGLYKNTCVGSDNVTDFNAIATCDWTNAGDYILANTCTERLKLFAAET LKATEETFKLSYGIATVREVISDRELHISWEVGKPRPPINRNYYFTGYRVTKNSKVOI GEYTFEKGDYGDAVVYRGTTTYKLNVGDYFVLTSHTVMPLSAPTLVPQEHYVRITGLY **AYVENTSITIKKPNELSLALGLKTIATHGIAAINSVPWSKILAYVKPFLGQAAITTSN** HTLSIHDGYTNEIIGYKAIQDGVTRDIISTDDCFANKHAGFDAWFSQRGGSYKNDKS(I SAKNRARTVAGVSI CSTMTNRQFHQKLLKSI AATRGATVVI GTSKFYGGWHNMLKTV RPFLCCKCCYDHVISTSHKLVLSVNPYVCNAPGCDVTDVTQLYLGGMSYYCKSHKPPI PTLNISDEFSSNVANYQKVGMQKYSTLQGPPGTGKSHFAIGLALYYPSARIVYTACSH SAINRPQIGVVREFLTRNPAWRKAVFISPYNSQNAVASKILGLPTQTVDSSQGSEYDY AENVTGLFKDCSKI ITGLHPTQAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMN GFKMNYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVN LVAVPTGYVDTENNTEFTRVNAKPPPGDOFKHLIPLMYKGLPWNVVRIKIVOMLSDTL FDYVYNPFMI DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVRRV DWSVEYPI I GDELRVNSACRKVQHMVVKSALLADKFPVLHDIGNPKAI KCVPQAEVEW

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LETLSRERVNINIVGDFHLNEEVALILASFSASTSAFIDTIKSLDYKSFKTIVESCGN
YVYKYKKPWYKOWNIGQDRSYLTPLCGEPSQAPAGNIRSIPARTLDAANHSIPDLQRAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NKALEMCI DOVTI ACAKLRSI.NI GEVPI AQSKGI.YRQCI RĞKEQLQI.LMPLKAPKEVT
FLEGDSHDTVLTSEEVVLKNGELEALETPVDS FTNGA I VGTPVCVNGLMI.LE I KDKBQ
YCALS PGILLATNNV FRI.KGGAPI KGVT FGEDTVWEVQGYKNVR I TFELDERVDKVI.NE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AKKCLHVVGPNLNAGEDĪQLLKĀAYENFNSQĪILLAPLLSAGIFGAKPLĢSLQVCVQT
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DFLKATCEHCGTENLVI EGPTTCGYL PTNAVVKMPCPACQDPE I GPEHSVADYHNHSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VGDVITSGDITCVVIPSKKAGGTTEMLSRALKKVPVDEYITTYPGQGCAGYTLEEAKT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NTNLHTQLVDMSMTYGQQFGPTYLDGADVTKIKPHVNHEGKTFFVLPSDDTLRSEAFE
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ch 90.9%; Score 20; DB 14; Length 29573; 1 Similarity 75.0%; Pred. No. 6.5; 15; Conservative 5; Mismatches 0; Indels 0
                                                             1 GUGAACUCACUCGUGAGCUC 20
 Query Match
                 Local
                 Best Loca
Matches
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RNA
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|:||||:|||:|||
|GTGAACTCACTCGTGAGCTC 726
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AY338175
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VRL 28-JUL-2003
                                                                                                  SARS coronavirus Taiwan TC2
SARS coronavirus Taiwan TC2
SARS coronavirus Taiwan TC2
Viruses; saRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
1 (Dases 1 to 29573)
Chang,J., Lin,T., Chen,C., Lin,T., Chan,W. and Shih,M.
linear
        SARS coronavirus Taiwan TC2, complete genome.
AY338175
AY338175.1 GI:32493130
                                                                              KEYWORDS
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ORGANISM
                                        ACCESSION
VERSION
                                                                                                                                                                                       REFERENCE
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NPMVIVNAANIHLKHGGGVAGALNKATNGAMQKESDDYIKLNGPLTVGGSCLLSGHNL
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KPKI KACI DEVTTTLEETKFLTNKLLLFAD I NGKLYHDSQNMLRGEDMSFLEKDAPYM
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ALKKCKSAFYVLPSEAPNAKEEILGTVSWNLREMLAHAEETRKLMPICMDVRAIMATI
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QRTELGVEFLKRGDKIVYHTLESPVEFHLDGEVLSLDKLKSLLSIREVKTIKVFTTVD
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VVCKHCGQKTTTLTGVEAVMYMGTLSYDNLKTGVSIPCVCGRDATQYLVQQESSFVMM
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SFDNFKLTCSNTKFADDLNQMTGFTKPASRELSVTFFPDLNGDVVAIDYRHYSASFKK
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AYVENTSITIKKPNELSLALGLKTIATHGIAAINSVPWSKILAYVKPFLGQAAITTSN
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LAKGVALDGVLSTFVSAARQGVVDTDVDTKDVIECLKLSHHSDLEVTGDSCNNFMLTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WNTKHGSGALRELTRELNGGAVTRYVDNNFCGPDGYPLDCIKDFLARAGKSMCTLSEQ
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                                                                                                                                                                              wmission (09-JUL-2003) Department of Molecular Medicine, China (09-JUL-2003) Taichung, Taichung, Taichung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (28-JUL-2003) Department of Molecular Medicine, China Medical University Hospital, 2, Yuh Der Road, Taichung, Taichung 404, Taiwan
                                                                                                                                                                                                                                                                   Medical University Hospital, 2, Yuh Der Road, Taichung,
                                                                                              2 (bases 1 to 29573)
Changy.J., Lin,T., Chen,C., Lin,T., Chan,W. and Shih,M.
Direct Submission
Submitted (09-JUL-2003) Department of Molecular Medicii
                                                                                                                                                                                                                                                                                                                      404, Taiwan
3 (bases 1 to 29573)
Changy, Lin, T., Chen, C., Lin, T., Chan, W. and Shih, M.
Direct Submission
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                                                                                                                                                                                                                                                                        Viruses; SBRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
1 (bases 1 to 29573)
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2 (bases 1 to 29573)
2 (Chang, J. G., Lin, T.-H., Chen, C.-M., Lin, T.-S., Chan, W.-L. and Shih, M.-C.
Direct Submission
                                              Gaps
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                                             Indels
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SARS coronavirus Taiwan TC3, complete genome.
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0
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Pred. No. 6.5;
5; Mismatches 0;
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ilarity 75:0%;
Conservative
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                                                                                                                                                                                        DEFINITION
                                                                                                                                                                                                                                                                 ORGANISM
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VERSION :
KEYWORDS
SOURCE
                                                                                                                                                                                                                                                                                                                                                  TITLE
JOURNAL
REFERENCE
AUTHORS
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LOCUS
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TITLE

FEATURES

CDS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LIANIFTPLVQPVGALDVSASVVAGGIIAILVTCAAYYFMKFRRVFGEYNHVVAANAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LFLMSFT1LCLVPAYSFLPGVYSVFYLYLTFYFTNDVSFLAHLQWFAMFSP1VPFW1T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YCALSPGLLATNNVFRLKGGAPIKGYTFGEDTVWEVQCYKNVRITFELDERVDKVLNË
KCSVYTVESGTEVTEFACVVAEAVVKTLQPVSDLLTNMGIDLDEWSVATFYLFDDAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ASFYYIWKSYVHIMDGCTSSTCMMCYKRNRATRVECTTIVNGMKRSFYVYANGGRGFC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NKVENMTPRDLGACIDCNARHINAQVAKSHNVSLIWNVKDYMSLSEQLRKQIRSAAKK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TSACVLAAECTIFKDAMGKPVPYCYDTNLLEGSISYSELRPDTRYVLMDGSI1QFPNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               YLEGSVRVVTTFDAEYCRHGTCERSEVGICLSTSGRWVLNNEHYRALSGVFCGVDAMN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WNTKHGSGALRELTRELNGGAVTRYVDNNFCGPDGYPLDCIKDFLARAGKSMCTLSEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LNSKVKVIQPRVEKKKTEGFMGRIRSVYPVASPQECNNMHLSTLMKCNHCDEVSWQTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DFLKATCEHCGTENLVIEGPTTCGYLPTNAVVKMPCPACQDPEIGPEHSVADYHNHSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENFSSRMYCSFYPPDEEEEDDAECEEEEIDETCEHBYGTEDDYOGLPLEFGASAETVR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VEEEEEEDWLDDTTEQSEIEPEPEPTPEEPVNQFTGYLKLTDNVAIKCVDIVKEAQSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YFDKAGQKTYERHPLSHFVNLDNLRANNTKGSLPINVIVFDGKSKCDESASKSASVYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LAKGVALDGVLSTFVSAARQGVVDTDVDTKDVIECLKLSHHSDLEVTGDSCNNFMLTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NNI PFRLTCATTRQVVNVITTKI SLKGGKIVSTCFKLMLKATLLCVLAALVCYIVMPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HTLSIHDGYTNEIIGYKAIQDGVTRDIISTDDCFANKHAGFDAWFSQRGGSYKNDKSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LDYIESKRGVYCCRDHEHEIAWFTERSDKSYEHQTPFEIKSAKKFDTFKGECPKFVFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HLKNGTCGLVELEKGVLPQLEQPYVFIKRSDALSTNHGHKVVELVAEMDGIQYGRSG
Submitted (23-JUL-2003) Department of Molecular Medicine, China
Medical University Hospital, 2, Yuh Der Road, Taichung, Taichung
404, Taiwan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |oin(196. .13329,13329. .21416)
|note="ribosomal slippage"
                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     country="Taiwan"
                                                                                                                                                                                                                  Bource
                   JOURNAL
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CDS

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Gaps 0; Indels Score 20; DB 14; Pred. No. 6.5; ; Mismatches 0; 5; 90.9%; ilarity 75.0%; Conservative Query Match Best Local Similarity

COMMENT

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SARS coronavirus Sin846, complete genome. AY559094 GI:45645021 AY559094.1 AY559094 DEFINITION ORGANISM ACCESSION VERSION KEYWORDS SOURCE AY559094

Viruses; ssRNA positive-strand viruses, no DNA stage; Nidovirales; Coronaviridae; Coronavirus. 1 (bases 1 to 29577) SARS coronavirus Sin846 SARS coronavirus Sin846

Vega.V.B., Ruan.Y., Liu, J., Lee, W.H., Wei, C.L., Se-Thoe, S.Y., Tang, K.F., Zhang, T., Kolatkar, P.R., Ooi, E.E., Ling, A.E., Stanton, L.W., Long, P.M. and Liu, E.T.
Mutational dynamics of the SARS coronavirus in cell culture and human populations isolated in 2003
(er) BMC Infect. Dis. 4 (1), 32 (2004) REFERENCE AUTHORS

JOURNAL TITLE

REFERENCE AUTHORS PUBMED

Disease 1 to 29577)
Vega,V.B., Ruan,Y., Liu,J., Lee,W.H., Wei,C.L., Se Thoe,S.Y., Yaga,V.B., Zhang,Y.F., Liu,S., Kolatkor,P., Ooi,E.B., Ee,L.A., Etanton,L.W., Long,P.M. and Liu,E.T.
Direct Submission
Submitted (24-FEB-2004) Genome Institute of Singapore, 60, Biopolis Street #02-01, Genome, Singapore 1386782, Singapore
Location/Qualifiers TITLE JOURNAL

FEATURES

Gaps ö DB 14; Length 29577; Indels 1. .29577 /organism="SARS coronavirus Sin846" /mol_type="genomic RNA" /db xref="taxon:267396" /country="Singapore" ; 0 90.9%; Score 20; DB ilarity 75.0%; Pred. No. 6.5; Conservative 5; Mismatches 1 GUGAACUCACUCGUGAGCUC 20 |:||||:|||:|||:|||:| 50 GTGAACTCACTCGTGAGCTC 779 Best Local Similarity Matches 15; Conserv 760 Query Match Bource ORIGIN g ð

Viruses; seRNA positive-strand viruses, no DNA stage; Nidovirales; Coronavirides; Coronavirus.

I (bases 1 to 29592)

Yuan, Z., Zhang, X., Hu, Y., Lan, S., Zhou, Z., Wang, H. and Wen, Y. Analysis of SARS coronavirus genome in Shanghai isolates

Lupublished

Z (bases 1 to 29592)

Yuan, Z., Zhang, X., Hu, Y., Lan, S., Zhou, Z., Wang, H. and Wen, Y. Direct Submission

Nimetred (11-NOV-2003) Key Lab of Medical Molecular Virology, Shanghai Medical College, Fudan University, 138 Yi Xue Yuan Road, Shanghai 200032, P.R. China

On or before Jan 5, 2004 this sequence version replaced gi:32454355, gi:32454355, gi:32454356, VRL 05-JAN-2004 SARS coronavirus ShanghaiOXCI, complete genome. AY463059 AH013000 AY322197 AY322198 AY322199 AY463059.1 GI:40457433 29592 bp SARS coronavirus ShanghaiQXC1 SARS coronavirus ShanghaiQXC1 AY463059 KEYWORDS SOURCE ORGANISM DEFINITION ACCESSION VERSION AUTHORS TITLE TITLE JOURNAL RESULT 15 REFERENCE JOURNAL AUTHORS REFERENCE AY463059 LOCUS

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VRL 24-SEP-2004

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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- nucleic search, using sw model OM nucleic

June 23, 2005, 23:06:41 ; Search time 192.283 Seconds (without alignments) 677.304 Million cell updates/sec Run on:

US-10-848-737-1 22

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Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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16.8 16.8 16.8	16.8	16.8	16.8 16.8 16.8	16.8 16.8 16.8	4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4
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ALIGNMENTS

small interfering RNA; siRNA; modified ribonucleotide, viral replication inhibition; hepatitis C virus; HCV; hepatitis C; antiinflammatory; hepatotropic; virucide; hepatitis A virus; hepatitis D virus; hepatitis E virus; Ebola virus; influenza virus; rotavirus; reovirus; poliovirus; human papilloma virus; metapneumoniavirus; coronavirus; viral infection; gene; ds. SARS coronavirus nucleotide sequence. BP ADJ39000 standard; DNA; 29751 25-JUL-2003; 2003WO-US023104. 26-JUL-2002; 2002US-0398605P (first entry) SARS coronavirus. #O2004011647-A1 05-FEB-2004. 06-MAY-2004 ADJ39000; RESULT 1 ADJ39000

(CHIR) CHIRON CORP.

Houghton M; Seo MY, Han J,

WPI; 2004-143862/14.

New RNase resistant small interfering RNA, useful for treating viral infections, e.g., hepatitis C, influenza virus or coronavirus infection.

Example 10; Fig 3; 74pp; English.

The present invention describes a small interfering RNA (siRNA) which comprises a modified ribonucleotide, where the siRNA is resistant to RNase and retains the ability to inhibit viral replication. Also described: (1) inactivating a virus in a patient, (2) making a modified siRNA that targets a nucleic acid sequence in a virus; (3) a double-

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         comprising a DNA molecule of 10-30 nucleotides that inhibits replication of hepatitis C virus (HCV); (4) inducing targeted RNA interference toward HCV; (b) treating he RNA molecule; (7) a host cell comprising a DNA segment encoding the RNA molecule; (7) a host cell comprising the vector of (6); (8) inhibiting replication of HCV in cells carrying HCV; (9) treating hepatitis C in a subject; (10) a modified siRNA molecule comprising a double-stranded RNA molecule of 10-30 nucleotides in length, which mediates RNA interference toward a target agent or virus and is linked to at least one receptor-binding ligand; and coll inducing targeted RNA interference in a patient. The modified siRNA molecules have antiinflammatory, hepatotropic and virucide activities. The modified RNA molecules are useful for incertivating virus in mammalian cells. The siRNAs are useful for treating hepatitis C virus, hepatitis D virus, hepatitis E virus, BDola virus, influenza virus, netapneumoniavirus or correct or compensate for cellular physiological abnormalities involved in conferring susceptibility to viral infections in patients and/or alleviate symptoms of a viral infection in patients. The present sequence represents the SRAS coronavirus nucleotide sequence, which is used in an example from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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secreted protein-related disease; transgenic animal, drug screening;
pharmacogenomic analysis; single nucleotide polymorphism; SNP.
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/standard name= "Single nucleotide polymorphism"
replace(8493, T)
/*tag= h
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/*replace(6100, T)
/*re
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replace(6122, A)
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replace(5479, G)
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replace(7641, G)
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replace(6883, G)
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                                                                                                                                                                                                                                                                                                                                                          90.9%; Score 20; DB 12; Length 29751; 75.0%; Pred. No. 8;
                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted protein genomic DNA coding sequence.
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Standard_name= "Single nucleotide polymorphism"
eplace(24159, G)
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standard_name= "Single nucleotide polymorphism"
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|standard_name= "Single nucleotide polymorphism"
|eplace(26339, A)
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'standard name= "Single nucleotide polymorphism"
:eplace(25508, G)
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standard name= "Single nucleotide polymorphism"
eplace(26693, A)
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standard name= "Single nucleotide polymorphism"
eplace(26884, T)
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standard_name= "Single nucleotide polymorphism"
eplace(28599, A)
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\fracepage bd name= "Single nucleotide polymorphism"
replace(35944, G)
                                   name= "Single nucleotide polymorphism"
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'etandard_name= "Single nucleotide polymorphism"
'eplace(23118, C)
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standard_name= "Single nucleotide polymorphism"
eplace(27320, T)
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|standard name= "Single nucleotide polymorphism"
|eplace(27339, A)
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eplace(27542, G)
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standard name= "Single nucleotide polymorphism"
eplace(28591, T)
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standard_name= "Single nucleotide polymorphism"
eplace(30857, A)
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standard_name= "Single nucleotide polymorphism"
eplace(31503, T)
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|seplace(22248, G)
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eplace(26582, G)
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standard_name= "Single nucleotide polymorphism"
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/standard_name= ":
replace(23789, A)
                                                    replace (22023, G)
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 replace(20491, T)
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                                                                                           /standard_name= "Single nucleotide polymorphism"
replace(39242, G)
/*tag= bh
/standard_name= "Single nucleotide polymorphism"
replace(39404, A)
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/standard name= "Single nucleotide polymorphism"
replace(40395, C)
/*tag= bj
                                                    /standard_name= "Single nucleotide polymorphism"
replace (37603, T)
/*tag= bg
/standard_name= "Single nucleotide polymorphism"
"Single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                 Length 340449;
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                                                                                                                                                                                                                                                                                                 DB 8;
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                                                                                                                                                                                                                                                                                               Score 18.8; DB
Pred. No. 48;
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                267426 GTGAACTCTCTCCTGAGCTCTT 267447
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 /standard_name= "
replace(37157, G)
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2001US-0303810P.
2001US-0315047P.
2001US-0324928P.
2001US-0330462P.
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2002US-0357844P.
2002US-0364134P.
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2001US-0336144P,
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68.2%;
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                                    *tag= bf
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                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 68.2
Matches 15; Conservative
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-OCT-2001
-NOV-2001
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1-FEB-2002
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Matches
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REPRESENTATION OF THE PROPERTY OF THE PROPERTY
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                                                                                                                                                                                                                          The invention relates to a novel method of predicting at least one toxic effect of a compound. The method comprises a gene expression profile of a tissue or cell sample exposed to the compound, and comparing the gene expression profile to a database comprising at least part of the data or information given in the specification. The methods are useful for predicting at least one toxic effect of a compound, predicting the progression of a toxic effect of a compound, predicting the progression of a toxic effect of a compound, predicting the renal toxicity of a compound, or identifying toxicity markers in tissues or cells exposed to known renal toxin. The genes are useful as toxicity markers in drug screening and toxicity assays, in monitoring disease or physiological states, or disease progression. This polynucleotide represents a rat DNA sequence relating to the toxic effect database described in the specification. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the World Intellectual Property
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                                                              Predicting at least one toxic effect of a compound, useful for toxicity modeling, comprises preparing a gene expression profile of a tissue or cell sample exposed to the compound, and comparing the gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        78.2%; Score 17.2; DB 10; Length 4113; 63.6%; Pred. No. 1.6e+02; ive 5; Mismatches 3; Indels 0;
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                                                                                                                                                                                     Example 4; Page; 446pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENE-) GENE LOGIC INC.
                                                                                                                                        profile to a database.
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                   WPI; 2003-148464/14.
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Elashoff M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Organization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2004.
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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The invention relates to a method of predicting (the progression of) a toxic effect of a compound by preparing a gene expression profile of a chidney tissue or cell sample exposed to the compound and comparing the gene expression profile to a database, or detecting the level of gene (s) expression in a tissue or cell sample exposed to the compound, where differential gene expression compared to a control indicates a toxic effect (toxicity progression). The method is useful for predicting (the progression of) at least one toxic effect of a compound. The genes are useful as toxicity markers in drug screening and toxicity assays. The methods are useful for predicting the likelihood that a compound or test agent will induce various specific kidney pathologies, such as nephritis, kidney necrosis, glomerular and tubular injury, or focal segmental capmental screening are useful for determining the similarity of a toxic response to one or more individual compounds and for predicting or endelling the potential cellular pathways influenced, induced or modulated by the compound or test agent. The kit is useful for predicting or modelling the potential cellular pathways influenced, induced or modulated by the compound or test agent. The kit is useful for monitoring the progression of renal disease states, for identifying genes that show promise as new drug targets and for screening known and newly designed drugs. This sequence corresponds to a gene marker used in the method of the invention. (Note: The sequence dor modelling prediction, but was obtained in electronic form part of the printed specification, but was obtained in electronic form part of the printed specification, but was obtained in electronic form part of the printed specification, but was obtained in electronic form and electronic form part of the printed specification, but was obtained but because in the sequence of the propression of the form will be sequence as the sequence of the propression of the sequence as the sequence of the sequence as the sequence
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gene expression profile of a kidney tissue or cell sample exposed to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.2%; Score 17.2; DB 12; Length 4113; 63.6%; Pred. No. 1.6e+02; ive 5; Mismatches 3; Indels 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4113 BP; 970 A; 1170 C; 1052 G; 921 T; 0 U; 0 Other;
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/product= "pRakCaR 3A 4Kb fragment"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat kidney calcium receptor 3A gene 4Kb fragment.
                                                                                                                                    Claim 11; SEQ ID NO 1417; 266pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV26965 standard; cDNA to mRNA; 4131 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3950 GTGAACTGACTGGTGTCTT 3971
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92US-00834044.
92US-00934161.
93US-00017127.
93US-00141248.
94US-00292827.
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Les 14; Conservative
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19-AUG-1994;
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21-AUG-1992
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                                                         compound
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The present sequence encodes rat kidney cell calcium receptor 3A (RakCaR 3A). The apecification includes details of molecules that can modulate one or more inorganic ion receptor activities, and antibodies and antibody fragments targetted to inorganic ion receptor proteins. The proteins, nucleic acids and antibodies may be used to treat disorders by modulating one or more inorganic ion receptor activities, preferably disorders of calcium homeostasis, e.g. hyperparathyroidism and
                                                                                                                                                                                  DNA encoding calcium receptor polypeptide(s) - useful for therapeutic purposes, e.g. hyperparathyroidism and osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parathyroid; calcium receptor; inorganic ion receptor; modulator; receptor expression; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 78.2%; Score 17.2; DB 2; Length 4131; Best Local Similarity 63.6%; Pred. No. 1.6e+02; Matches 14; Conservative 5; Mismatches 3; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4131 BP; 987 A; 1170 C; 1053 G; 921 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat parathyroid calcium receptor 3A nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/product= "RakCaR 3A"
/note= "parathyroid calcium receptor"
                                                                                             Hebert SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ25056 standard; cDNA to mRNA; 4131 BP.
                                                                                                                                                                                                                                     Claim 15; Col 133-142; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3950 GTGAACTGACTGGTGTGCTCTT 3971
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574. .3813
                                                                                             Brown EM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NPSP-) NPS PHARM INC.
(BGHM ) BRIGHAM & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Garrett JE;
                                        (BGHM ) BRIGHAM & WOMENS HOSPITAL (NPSP-) NPS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GUGAACUCACUCGUGAGCUCTT
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93US-00141248.
94US-00292827.
94WO-US012117.
     94US-00353784.
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                                                                                           Garrett JE, Fuller FH,
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                                                                                                                              WPI; 1998-008040/01.
P-PSDB; AAW38275.
       08-DEC-1994;
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08-DEC-1994;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                     osteoporosis
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22-OCT-1993;
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                                                                                                                                                                                                                                                                                                      The tissue from which this receptor and receptors from bovine parathyroid and rat kidney are derived, respond to changes, and control changes, in calcium ion concentration, e.g. parathyroid hormone regulates Ca2+ homeostasis in blood and extracellular fluid, and kidney function alters through changes in Ca2+ levels in juxtaglomerular and proximal tubule cells in the kidney. The purified receptors (produced recombinantly) can be used to screen for compounds that modulate calcium receptor activity, especially those that can be used to treat diseases associated with the receptors in these tissues. They can also be used to raise antibodies for
                                                                                                                                                                                                 Calcium receptor poly:peptide(s) - useful for drug screening or antibody
                                                                                                                                                                                                                                                                                         The rat kidney calcium receptor gene encodes a 1079 amino acid protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          calcium receptor 3A; RakCaR 3A; calcium homeostasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78.2%; Score 17.2; DB 2; Length 4131; 63.6%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4131 BP; 988 A; 1170 C; 1052 G; 921 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/product= "kidney_cell_calcium_receptor_3A"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rat kidney cell calcium receptor 3A (RakCaR 3A) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT95860 standard; cDNA to mRNA; 4131 BP
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574. .3813
                                                                                                                                                                                                                                                  Example 30; Fig 50; 174pp; English.
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                                                                        (BGHM ) BRIGHAM & WOMENS HOSPITAL.
                                                                                                            Garrett JE;
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92US-00834044.
92US-00934161.
93US-00009389.
93US-00141248.
94WG-00292827.
 94WO-US012117.
94US-00353784.
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Best Local Similarity 63.6'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use in detection assays
                                                    (NPSP-) NPS PHARM INC.
                                                                                                          Hebert SC, Brown EM,
                                                                                                                                           WPI; 1998-347412/30.
P-PSDB; AAW54847.
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21-OCT-1994;
08-DEC-1994;
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19-AUG-1994;
21-OCT-1994;
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AAT95860

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94US-00353784.

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08-DEC-1994;
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                                                                                                                            3A). Calcium receptor polynucleotides may be used: (i) for producing activity velationships, for assaying molecular activity on the receptor (i.e. identifying modulators of receptor function) and for producing antibodies specific for the receptor function) and for producing antibodies specific for the receptor; (ii) for sequencing the normal form of the nucleic acids (the derived sequence may be compared with other receptors to identify conserved sequences, mutations and variations that may influence calcium receptor activity and to determine target sites for antisense molecules ribozymes, hybridization probes and polymerase chain reaction (PCR) amplification primers); (iii) as hybridization probes to detect the presence of similar sequences in samples; and (iv) as PCR primers to generate particular nucleic acid sequence regions; (e.g. to generate regions to be probes by hybridization detection probes)
                                            Nucleic acids encoding protein calcium receptors useful for identifying modulators of receptor expression and activity and for the production of antigens specific for calcium receptors.
                                                                                                                   The present sequence encodes rat parathyroid calcium receptor 3A (RakCaR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parathyroid calcium receptor; inorganic ion receptor; osteoporosis; calcium homeostasis; hyperparathyroidism; seizure; stroke; epilepsy; spinal cord injury; hypoxia-induced nerve cell damage; cardiac arrest; neonatal distress; neurodegenerative disease; Alzheimer's disease; Huntington's disease; Parkinson's disease; dementia; muscle tension;
                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                     78.2%; Score 17.2; DB 2; Length 4131; 63.6%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                 Sequence 4131 BP; 988 A; 1170 C; 1052 G; 921 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat parathyroid calcium receptor pRakCaR 3A encoding cDNA.
                                                                                                                                                                                                                                                                                                                                               5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV82486 standard; cDNA to mRNA; 4131 BP.
                                                                                                                                                                                                                                                                                                                                                                                    3950 GTGAACTGACTGGTGTGCTCTT 3971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                      1 GUGAACUCACUCGUGAGCUCTT 22
                                                                                            Claim 8; Fig 50; 174pp; English.
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92US-00934161.
93US-00017127.
93US-00141248.
94WS-00292827.
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/*tag= a
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Best Local Similarity 63.6
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               depression; anxiety; ss
           WPI; 1999-571274/48.
P-PSDB; AAY41781.
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22-OCT-1993;
19-AUG-1994;
21-OCT-1994;
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21-AUG-1992
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A method has been developed of screening for a compound able to affect one or more activities of a calcium receptor (CR) comprises: (A) contacting a recombinant cell with a test compound, where the recombinant cell comprises a recombinant mucleic acid expressing the CR, provided that the cell does not have functional CR expression from endogenous nucleic acid; (B) determining the ability of the test compound to affect one or more activities of the calcium receptor; and (C) comparing the ability with the ability of the test compound to affect the one or more activities in a cell not comprising the recombinant nucleic acid. The present sequence encodes rat parathyroid CR, designated a pRakCaR 3A. The nucleic acid sequence of pRakCaR 3A can be used as part of the recombinant nucleic acid in the method described above. The compounds identified can be used to treat diseases or disorders characterised by enhormal calcium homeostasis, e.g. spyperparathyroidism, osteoporosis and other bone and mineral-related disorders. They can also be used for the treatment of diseases and disorders. They can also be used for the treatment of diseases and disorders associated with disrupted Ca2+ responses, e.g. seizures, stroke, spinal cord injury, hypoxia-induced nerve cell damage such as in cardiac arrest or neonatal distress, epilepsy, neurodegenerative diseases such as Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                Screening for calcium receptor-active compounds - by recombinant expression of nucleic acid encoding calcium receptor and determining the effect of compounds on calcium receptor activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Huntington's disease and Parkinson's disease, dementia, muscle tension,
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                                                                                                                           Brown EM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4131 BP; 988 A; 1170 C; 1052 G; 921 T; 0 U; 0 Other;
                                                                                                                      Nemeth EF,
                                                                                                                           Van Wagenen BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ89299 standard; cDNA to mRNA; 4131 BP.
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(NPSP-) NPS PHARM INC.
(BGHM ) BRIGHAM & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 50; 176pp; English.
                                                                                                                      Balandrin MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               574. .3813
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les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 depression, and anxiety
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                                                                                                                                                                 Hebert SC;
                                                                                                                                                                                                                                             WPI; 1999-119871/10.
P-PSDB; AAW89566.
                                                                                                                      Del Mar EG,
Garrett JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rattus sp.
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AAI72123;

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This invention describes a novel method of treating disorders by

calcimimetic, or a calcilytic compound. The products of the invention

calcimimetic, or a calcilytic compound. The products of the invention

have osteopathic, cerebroprotective, cytostatic, neuroprotective,

dermatological, tranquilizer, vulnerary, antiulcer, immunosuppressive,

thypotensive and cardiant activity. The method is suitable for reducing

parathyroid hormone level in a patient to that of a normal individual,

treating a patient having osteoporosis, to inhibit bone resorption, and

creating a patient having osteoporosis, to inhibit bone resorption, and

creating a patient having osteoporosis, to inhibit bone resorption, and

creating a patient having osteoporosis, to inhibit bone resorption, and

creating a patient having osteoporosis, to inhibit bone resorption, and

creating a patient having osteoporosis, as decrease in plasma Ca2+. The

method is useful in treating disorders in humans such as

method is useful in treating disorder in vivo. The level of

creating and patient having osteoporosis. Also for treatment

creating a patient having osteoporosis, product or prevention, based on the affected cells, of other disorder,

cronditions like seizures, stroke, head trauma, spinal cord injury,

hypoxia-induced nerve cell damage such as in cardiac arrest or neonatal

creating aliesase, dementia, muscle tension, depression, anxiety, panic

disorder, OCD (not defined), post-traumatic stress disorder,

creat inappropriate ADH serretion (SIDH), cirrhosis, congestive

creat failure and nephrosis, hypertension, preventing and/or decreasing

creation of inappropriate ADH serretion (SIDH), cirrhosis, congestive

creation of inappropriate ADH serretion (SIDH), cirrhosis, condensing

creations, gut motility disorders such as diarrhoes and spastic colon,

creations, and services and antibiotics (e.g. aminoglycoside

antibiotics), gut motility disorders such as diarrhoes and spastic colon,

creations and appropriate ADH servetone of antibiotics of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating disorders or diseases in a patient by modulating inorganic ion receptor activities especially calcium receptor for hyperparathyroidism, by administering a calcimimetic or calcilytic compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rat calcium receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Van Wagenen BC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nemeth EF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BGHM ) BRIGHAM & WOMENS HOSPITAL.
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92US-00934161
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93US-00009389
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94US-00292827
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P-PSDB; AAY51828.
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                                                                                                                                                                        07-JUN-1995;
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US6031003-A
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21-AUG-1992
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The sequences given in AA172120-23 encode various calcium receptor
proteins. This sequence encodes a rat calcium receptor. The calcium
receptor proteins are used, in conjunction with the compounds of the
invention, for structure determination, to assay a molecules activity on
a receptor and to obtain antibodies to that receptor. The compounds of
the invention, which modulate inorganic ion receptors are useful for
treating and diagnosing diseases or disorders which can be treated by
modulating inorganic ion receptor activity. This is preferably a disease
or disorder characterized by abnormal inorganic ion homeostasis,
cor disorder characterized by abnormal inorganic ion homeostasis,
steeporosis and other bone and mineral-related disorders), an abnormal
level of a messenger whose production or secretion is affected by
inorganic ion receptor activity, and an abnormal level or activity of a
messenger whose function is affected by inorganic ion receptor activity.
These compounds are also useful for treating and diagnosing diseases or
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                                                                                                                                                                                                                                       Gaps
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                                                                                                                                        Score 17.2; DB 3; Length 4131;
Pred. No. 1.6e+02;
5; Mismatches 3; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                  3950 GTGAACTGACTGGTGTCTCTT 3971
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                                                                                                                                        Query Match
Best Local Similarity 63.6%;
Matches 14; Conservative
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AAI72123 standard; cDNA to mRNA; 4131 BP.

RESULT 10 AAI 72123

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c-cell; inorganic ion receptor; homeostasis; sarcoidosis; hyperparathyroidism; osteoporosis; central nervous system; seizure; stroke; head trauma; spinal cord injury; organ transplant rejection; hypoxia-induced nerve cell damage; cardiac arrest; neonatal distress; epilepsy; neurodegenerative disease; Alzheimer's disease; cirthosis; huntington's disease; parkinson's disease; dementia; depression; anxiety; panic disorder; obsessive-compulsive disorder; spatic colon; post-traumatic stress disorder; schizophrenia; diarrhoea; kidney; neuroleptic mallapant syndrome; Tourette's syndrome; gut motility; inappropriate ADH secretion; SIADH; gastrointestinal ulcer disease; congestive heart failure; nephrosis; hypertension; alternative splicing;
                                                                                     Calcium receptor; bovine; human; parathyroid; calcium receptor; thyroid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel inorganic ion receptor-modulating compounds, useful for treating e.g. hyperparathyroidism, osteoporosis, stroke, epilepsy, Alzheimer's disease, dementia, depression, anxiety, hypertension, cirrhosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Delmar EG, Nemeth EF;
                                                                                                                                                                                                                                                                                                                                    /product= "RakCaR3A"
                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                             aminoglycoside antibiotic; ss.
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92US-00934161.
93US-00017127.
93US-00141248.
94US-0012827.
94US-00353784.
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                                    25-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                           574. .3813
                                                            cDNA encoding pRakCaR3A
                                                                                                                                                                                                                                                                                                                           /*tag=
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P-PSDB; AAB47823.
                                                                                                                                                                                                                                                                       Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      spastic colon
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Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; adrenal disorder;

Transgenic disorder; blood disorder; immune disorder; bone disorder;

Transgenic disorder; metabolic disorder; nutritive disorder; cancer;

Transgenic disorder; metabolic disorder; nutritive disorder; cancer;

Transgenic disorder; prostate disorder; breast disorder;

Transgenic disorder; thyroid disorder; pancreas disorder; spleen disorder;

Try disorder; thyroid disorder; nutrianginal; antimaric;

Try cytostatic; antihilammatory; vasotropic; antianginal; antimarkythmic;

Try dermatological; antibulcer; antibyroid; antiallargic; antiseborrhoeic;

Try dermatological; antibulcer; antibyroid; antiallargic; anorectic;

Try mmunosuppressive; nephrotropic; gene therapy; GPCR modulator; mouse;
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disorders of the central nervous system such as seizures, stroke, head trauma, spinal cord injury, hypoxia-induced nerve cell damage such as in cardiac arrest or neonatal distress, epilepsy, neurodegenerative diseases such as Alzheimer's disease, luntington's disease and Parkinson's disease, dementia, depression, anxiety, panic disorder, obsessive-compulsive disorder, post-traumatic stress disorder, obsessiveneuroleptic malignant syndrome and Tourette's syndrome, diseases involving excess water reabsorption by the kidney such as inappropriate ADH secretion (SIADH), cirrhosis, congestive heart failure, nephrosis, hypertension, for preventing and/or decreasing renal toxicity from cationic antibiotics (e.g. aminoglycoside antibiotics), gut motility disorders such as diarrhoea, and spastic colon, gastrointestinal (GI) ulcer diseases, GI diseases with excessive calcium absorption such as sarcoidosis, and autoimmune diseases and organ transplant rejection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     78.2%; Score 17.2; DB 6; Length 4131; 63.6%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4131 BP; 988 A; 1170 C; 1052 G; 921 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hohmann J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaitanaris GA, Bergmann JE, Gragerov A, Hohmann J,
Madisen L, Mcilwain KL, Pavlova MN, Vassilatis D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse GPCR CASR polynucleotide, SEQ ID NO:1217.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1.6e
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3950 GTGAACTGACTGGTGTGTTT 3971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GUGAACUCACUCGUGAGCUCTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADO30115 standard; cDNA; 4550 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-2003; 2003WO-US028226.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-SEP-2002; 2002US-0409303P,
09-APR-2003; 2003US-0461329P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 63.6
les 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PRIM-) PRIMAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            murine; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB; ADO29212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004040000-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-MAY-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO30115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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CC (GPCRs) and nucleic acids encoding them. The invention also relates to sequences at least 90% identical to the GPCR proteins and nucleic acids encoding them. The invention also relates to sequences at least 90% identical to the GPCR proteins and nucleic acids of the invention; methods of treating, preventing or diagnosing diseases associated with GPCRs of the invention; methods of screening for compounds useful in the treatment of GPCR-related diseases; a transgenic mouse comprising a GPCR pene of the invention; a mouse comprising a GPCR pene of the invention; and with the transgene or in an endogenous GPCR gene; cells derived from the transgene or in an endogenous GPCR gene; cells derived from the transgenic mice; kits comprising several mice, each of which has a mutation in a different GPCR gene of the invention; and kits comprising probes which hybridise to GPCR polymucleotides of the invention. The invention intruther discloses variants of the GPCR polympetides and vectors comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may be used in the diagnosis, treatment or prevention of a wide variety of discorders of the adrenal gland; disorders (e.g., Alzhaimer's disease, diarrhoea, food poisoning or irritable bowel syndrome); cardiovascular disorders (e.g., Alzhaimer's disease, diarrhoea, food poisoning or irritable bowel syndrome); cardiovascular disorders (e.g., autoimmune disorders (e.g., anomania or laukaemia); immune disorders (e.g., autoimmune disorders (e.g., cancers); metabolic or nutritive disorders or carterities, gout or osteoporosis); metabolic or vitamin deficiency-related diseases or vitamin deficiency-related diseases); and disorders (e.g., autoimmune disorders (e.g., cancers); the present sequence represents a GPCR-encoding nucleic acid of the invention. Note: The full sequence deta for this prober tide not form part of the present sequence represents a GPCR-encoding content with one potationary electronic format directly formed to response to the sequence represents a GPCR-en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
associated with GPCR dysfunction e.g. autoimmune diseases, angina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4550 BP; 1107 A; 1287 C; 1126 G; 1030 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 78.2%; Score 17.2; DB 12; Local Similarity 63.6%; Pred. No. 1.6e+02; les 14; Conservative 5; Mismatches 3;
                                                                     Claim 151; SEQ ID NO 1217; 542pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3891 GTGAACTGACTGGTGGGCTCTT 3912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GUGAACUCACUCGUGAGCUCTT
                        pectoris, Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACN44600 standard; DNA; 57561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-2002; 2002US-00087192.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-FEB-2003; 2003WO-US006235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SAGR-) SAGRES DISCOVERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACN44600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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The first strand primer ZC2938 was prepared from bovine cDNA and used as a template cDNA for PCR reactions. ZC2938 (SEQ ID No 9) was synthesised from one time poly d(T)-selected bovine liver poly (A)+ RNA in two separate reactions, to assess the quality of both first and second strand synthesis. PCR was carried out by stendard methods using a set of degenerate oligonucleotides complementary to the N-terminus of a gamma-carboxylase protein. Amplified DNA was ligated into a plasmid and the plasmids were selected for a gamma-carboxylase insertion. See also AAQ21467-77 (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                           Gamma-carboxylase protein compsns. - used in recombinant prodn. of active vitamin-K dependent proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human calcitonin receptor; hCR; bone resorption disorders; detection; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA encoding human calcitonin receptor - used to develop drugs for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.4%; Score 16.8; DB 2; Length 44; 70.0%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer to amplify cDNA of T-47D beast carcinoma cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 44 BP; 10 A; 7 C; 7 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kuestner RE;
                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 55; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 GAACUCACUCGUGAGCUCTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAATTCACTAGTGAGCTCTT 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ59009 standard; DNA; 44 BP
                                                                           90US-00557220.
                                                                                                                90US-00557220.
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93US-00100887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 70.0 Matches 14; Conservative
                                                                                                                                                                       (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sheppard PO,
                                                                                                                                                                                                                                                      WPI; 1992-064951/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-135573/16.
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                                                                           23-JUL-1990;
                                                                                                                                  14-MAR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-AUG-1993;
                                                                                                                23-JUL-1990;
WO9201795-A.
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16-NOV-1994
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                                      06-FEB-1992
                                                                                                                                                                                                                3erkner KL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ59009;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABD32923_5/c
Continuation (6 of 8) of ABD32923 from base 500001 (Human cancer-associated genomic DNA WP Sequence split into 8 fragments LOCUS ABD32923 Accession Abd32923
WP ABD32923_1 100001 10000
WP ABD32923_2 200001 310000
WP ABD32923_3 300001 410000
WP ABD32923_4 400001 510000
WP ABD32923_5 500001 610000
WP ABD32923_6 500001 610000
                                                                                                                                              The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (1) for screening drug candidates; (11) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (111) for screening of a bioactive agent capable of modulating the activity of CAP; (1v) for carcinoma; (vi) for inhibiting the activity of CAP; (vi) for diagnosing carcinoma; (vi) for inhibiting the effect of CAP; (ix) as a biochip; (x) for diagnosing carcinoma; (vii) for neutralizing the effect of CAP; (ix) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma associated (CA) gene copy number. In addition, the CAP genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent CA coding sequence. Note: This patent is an equivalent to basic patent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ô
                                                     Recombinant nucleic acid useful for diagnosis and treatment of carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 57561 BP; 14226 A; 12066 C; 13347 G; 16427 T; 0 U; 1495 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.2; DB 13; Length 110000;
Pred. No. 2.6e+02;
3; Mismatches 3; Indels 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78.2%; Score 17.2; DB 11; Length 57561; 63.6%; Pred. No. 2.3e+02; ive 5; Mismatches 3; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vitamin K dependent proteins; amplification; degenerate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZC2938- PCR primer for gamma-carboxylase proteins.
                                                                                                              Claim 1; SEQ ID NO 1129; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15732 GAGAACTCAGCCGTGAGCTCTT 15711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GUGAACUCACUCGUGAGCUCTT 22
                                                                         comprises a nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ21468 standard; cDNA; 44 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.2%;
72.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 63.57
These 14; Conservative
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Best Local Similarity 72.7
Matches 16; Conservative
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                WPI; 2003-328604/31.
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02-JUN-1992
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RESULT 14 AAQ21468

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Gaps

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PT treatment and prevention of bone resorption disorders and for detection XX XX PS Example 1; Page 61; 83pp; English.

XX CT-47D human breast carcinoma cells were used to prepare a cDNA library CC used for the cloning of calcitonin receptor sequences, AAQ59009 was used CC in first strand cDNA synthesis from the poly (A) + RNA. Human calcitonin CC receptors can be used in the development of drugs treatment of bone CC resorption disorders. (Updated on 25-MAR-2003 to correct PN field.)

XX Sequence 44 BP; 10 A; 7 C; 7 G; 20 T; 0 U; 0 Other;
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 Query Match
 76.4%; Score 16.8; DB 2; Length 44;

 Best Local Similarity 70.0%; Pred. No. 1.26+02;

 Matches 14; Conservative 4; Mismatches 2; Indels

 Qy 3 GARCUCACUCGAGGCUCTT 22

 QY | | | | | | | | | | | | | | | |

 Db 12 GAATTCACTAGTGAGCTCTT 31

0;

0; Gaps

Search completed: June 24, 2005, 00:22:06 Job time: 199.283 secs

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US-08-485-588-4
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65, Appl
12478, A
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                                                                           June 23, 2005, 23:58:22 ; Search time 56.4724 Seconds (without alignments) 637.445 Million cell updates/sec
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Sequence 4, 1
Sequence 4, 1
Sequence 38,
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Sequence 3
Sequence 3
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                                                                                                                                                                                                                                                                                                                                                                              Issued Patents NA:*

(cgn2_6/ptodata/1/ina/5A_COMB.seq:*

(cgn2_6/ptodata/1/ina/5B_COMB.seq:*

(cgn2_6/ptodata/1/ina/6A_COMB.seq:*

(cgn2_6/ptodata/1/ina/6B_COMB.seq:*

(cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

(cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

(cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-943-986-4
US-08-943-986-4
US-08-848-159-4
US-08-848-159-4
US-08-453-742-15
US-08-453-742-15
US-08-453-222-15
US-08-453-222-15
US-08-453-222-15
US-08-453-222-12
US-09-902-540-1000
US-09-902-540-1000
US-09-949-016-12214
US-09-949-016-12214
US-09-9433-110-1469
US-09-107-433-2198
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-09-949-016-12478
-09-949-016-15631
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US-09-949-016-13100
                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                               1202784 segs, 818138359 residues
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                                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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22
1 gugaacucacucgugagcuctt 22
                                                 - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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seq length: 2000000000
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24538
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FASTER
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,751
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/353,784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/08480751
Patent No. 5858684
GENERAL INFORMATION:
APPLICANT: Edward F. Nemeth
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 90071
COMPUTER READBLE FORM:
MEDIUMYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
APPLICATION NUMBER: U.S. 08/017,12/
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 13 August, 1991
ATTORNEY AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REGISTRATION NUMBER: 213/006
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEPX: 67-3510
INFORMATION FOR EGO ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 4131 base pairs
TYBE: nucleic acid
STRANDEDNESS: Single
TOWNER TOWNER TOWNER
THENDEN TOWNER
THENDER
THENDEN TOWNER
THENDE T
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; LOCATION: 574..3810
; OTHER INFORMATION:
US-08-484-565-4
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CITY: Los Angeles
STATE: California
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US-08-480-751-4
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,565
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA: 1995
BRIOR APPLICATION DATA: including application PRIOR APPLICATION DATA: described below: 9
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 23 Cotober, 1994
APPLICATION NUMBER: U.S. 08/09,389
FILING DATE: 23 Cotober, 1993
APPLICATION NUMBER: U.S. 08/099,389
FILING DATE: 23 February, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-44-565-4

Sequence 4, Application US/08484565

Sequence 4, Application US/08484565

Patent No. 5763569

GENERAL INFORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
TITLE OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 78.2%; Score 17.2; D
Best Local Similarity 63.6%; Pred. No. 47;
Matches 14; Conservative 5; Mismatches
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                                   FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/005
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (7-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 4131 base pairs
TYPE: nucleic acid
STRANDENDESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3950 GTGAACTGACTGGTGCTCTT 3971
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: CDNA to mRNA
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MEDIUM TYPE: 3.5" Di
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COTHER INFORMATION:

US-08-485-588-4
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Length 4131;
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APPLICANT: Brown, Steven C. Hebert,
APPLICANT: Bradford C. Van Wagenen, Manuel
APPLICANT: F. Balandrin, Forrest H. Fuller,
APPLICANT: Eric G. DelMar, and Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 17.2; DB Pred. No. 47; 5; Mismatches
                                                                                                      FILING DATE: 7-June-1995
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 OCCODER, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 OCCODER, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
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First Interstate World Center
Suite 4700
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 AUGUST, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Heber, Sheldon O. REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/006
TELECOMMUNICATION INFORMATION:
                                                                                          08/484,565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-353-784-4
; Sequence 4, Application US/08353784
; Patent No. 6011068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELERAX: (213,
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTER.STICS:
SAMPLE A 131 base pairs
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TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 63.6%
Marches 14; Conservative
                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: 574..3810
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OTHER INFORMATION
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APPLICANT: Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-943-986-4
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Patent No. 5962314
GENERAL HORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: Jewen E. Jewen E. TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Site 1000
STREET: First First First First Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Mismatches
                                                                                                                                          APPLICATION NUMBER: U.S. 08(009,389)
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08(017,127)
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 AUGUST, 1992
APPLICATION NUMBER: U.S. 07/934,044
FILING DATE: 21 AUGUST, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 AUGUST, 1991
ATTORNEY/AGENT INFORMATION:
                                      21 October, 1994
UMBER: U.S. 08/292,827
                                                                                    23 August, 1994
UMBER: U.S. 08/141,248
22 October, 1993
UMBER: U.S. 08/009,389
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9 December, 1994
JMBER: PCT/US/94
                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 213/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                       38,179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: cDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4131 base pairs
                                                                                                                                                                                                                                                                                                                                                                                             NAME: Heber, Sheldon O. REGISTRATION NUMBER: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 63.6'
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CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) LOCATION: 574.3810
; OTHER INFORMATION:
US-08-480-751-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Di
                                                           APPLICATION NUMBER:
FILING DATE: 23 Auguaphilication NUMBER:
FILING DATE: 22 Octo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
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STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nuclei
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-943-986-4
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DB 3; Length 4131;
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                                                                                                                                                                                                                                                    ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: MS Word
SOFTWARE: FastSEG for Windows Version 3.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,719B
FILING DATE: 7 June, 1995
                NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 78.2%; Score 17.2; Dest Local Similarity 63.6%; Pred. No. 47; Matches 14; Conservative 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 Aguat, 1994
APPLICATION NUMBER: U.S. 08/11,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/01,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 08/01,127
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/934,61
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/934,64
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/94,61
FILING DATE: 21 AUGUST, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 AUGUST, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3950 GTGAACTGACTGGTGCTCTT 3971
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Douglas C. Murdock
REGISTRATION NUMBER: 37,549
REFERENCE/DOCKET NUMBER: 213
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 4131 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION: 514
PRIOR APPLICATION NUMBER: 0
FILTING NAME FILTING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   574..3810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 7 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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US-08-484-719B-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ò
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                                                                                                                                                                                                                                      APPLICATION NUMBER:

FILING DATE: 9 December, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 8
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 Adust, 1993
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 Sebruary, 1993
APPLICATION NUMBER: U.S. 07/034,161
FILING DATE: 21 Aqust, 1992
APPLICATION NUMBER: U.S. 07/834,161
FILING DATE: 21 Aqust, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 21 Aqust, 1991
ATTORNEY/ABDAT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
APPLICATION NUMBER: 38,179
APPLICATION NUMBER: 38,179
APPLICATION NUMBER: 38,179
APPLICATION NUMBER: 38,179
TELEPHONE: (213) 499-1600
TELEPHONE: (213) 499-1600
TELEPHONE: (213) 499-1600
TELEPHONE: GLARACTERISTICS:
LENGTH: 4131 base pairs
TYPE: nucleic acid
TYPE: UNDER TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/08484719B
Patent No. 6031003
GENERAL INFORMATION:
APPLICANT: Edward F. Nemeth, Edward M.
APPLICANT: Brown, Steven C. Heibert,
APPLICANT: Bradford C. Van Wagenen,
APPLICANT: Manuel F. Balandrin,
APPLICANT: Forrest H. Fuller, Eric G.
APPLICANT: Forrest H. Fuller, Eric G.
APPLICANT: Delmar, Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: FASTSEQ CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/353,784
FILING DATE: 9 December, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       query Match 78.2%; Score 17.2; Best Local Similarity 63.6%; Pred. No. 47 Matches 14; Conservative 5; Mismatcher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3950 GTGAACTGACTGGTGCTCTT 3971
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: 574..3810
COTHER INFORMATION:
US-08-353-784-4
California
                                              90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 6
US-08-484-719B-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                      APPLICANT: Wooldike, Helle F.
APPLICANT: Hagen, Frederick
APPLICANT: Hjort, Carsten M.
APPLICANT: Sven, Hastrup
TITLE OF INVENTION: An Enzyme Capable of Degrading Cellulose
TITLE OF INVENTION: or Hemicellulose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CONFULAR: LEM FC COMPOSITION OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA: US/08/361,920 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Moore, Emma E
APPLICANT: Sheppard, Paul O
APPLICANT: Kuestner, Rolf E
TITLE OF INVENTION: Human Calcitonin Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.8; D
Pred. No. 43;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PELING DATE: 28-0CT-1992
APPLICATION NUMBER: US 07/940,860
FILING DATE: 09-MAY-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 1158/90
PRIOR APPLICATION DATA:
FILING DATE: 08-MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Elias J.
REGISTRACE/DOCKET NUMBER: 33,728
REFERRNCE/DOCKET NUMBER: 3435.204-US
TELECOMMUNICATION INFORMATION:
                     3950 GTGAACTGACTGGTGTCTCTT 3971
                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: New York
COUNTRY: United States of America
COUNTRY: United States of America
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                         US-08-361-920-38
; Sequence 38, Application US/08361920
; Patent No. 5457046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15, Application US/08453742
Patent No. 5622839
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 GAATTCACTAGTGAGCTCTT 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic) US-08-361-920-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 70.0%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 8:
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 54570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-453-742-15
                                                                                    RESULT 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/35,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: 10.5.08/292,827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      78.2%; Score 17.2; DB 3; 63.6%; Pred. No. 47;
APPLICANT: Eric G. Del Mar
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: LVON C. STRPET
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                              STREET: Lyon & Lyon STREET: First Interstate World Center STREET: Suite 4700 STREET: 633 West Fifth Street CITY: Los Angeles STATE: California COUNTRY: USA ZIP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 23 Aquast, 1994
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/011,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 12 Aquast, 1992
APPLICATION NUMBER: U.S. 07/844,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/844
APPLICATION NUMBER: U.S. 07/844
APPLICATION NUMBER: U.S. 07/844
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 Aquast, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,159
FILING DATE: 7 June, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REPERENCE/DOCKET NUMBER: 214/
TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 4131 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Heber, Sheldon O. REGISTRATION NUMBER: 38,179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 63.67
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: CDS
LOCATION: 574..381
COTHER INFORMATION:
US-08-484-159-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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APPLICANT: Sheppard, Paul O
APPLICANT: Sheppard, Paul O
APPLICANT: Kuestner, Rolf B
TITLE OF INVENTION: Human Calcitonin Receptor
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND AND TOWNSEND KHOURIE and CREW
STREET: One Market Plaza, Steuart St. Tower,
STREET: Twentieth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: END FC compatible
OPERATING SYSTEM: FC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/100,887
FILING DATE: 30-MAY-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/100,887
FILING DATE: 02-AUG-1993
APPLICATION NUMBER: US 07/954,804
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.4%; Score 16.8; I
70.0%; Pred. No. 43;
cive 4; Mismatches
FILING DATE: 30-MAY-1995
CIASSIFICATION: 435
FILING DATE: 30-MAY-1995
FILING DATE: 02-AUG-1993
FILING DATE: 02-AUG-1993
APPLICATION NUMBER: US 07/954,804
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W
REGISTATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 13952-15-1
FELECOMMUNICATION INFORMATION:
TELEFAX: 415-543-5043
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
TERRIDENCES: SINGLE
                                                                                                                                                                                                                                                 13952-15-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13952-15-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-453-222-15; Sequence 15, Application US/08453222; Patent No. 5674981; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Parmelee, Steven W
REGISTRATION NUMBER: 31,990
REPERENCE/DOCKET NUMBER: 1395;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-467-9600
TELEPAX: 415-543-5043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 GAACUCACUCGUGAGCUCTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 70.0
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear; IMMEDIATE SOURCE: CLONE: ZC2938 US-08-454-464-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94105-1492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-454-464-15
; Sequence 15, Application US/08454464
; Patent No. 567468
; GENERAL INFORMATION:
    APPLICANT: Moore, Emma E
    APPLICANT: Sheppard, Paul O
    APPLICANT: San KHOURIE and CREW
    STREET: One Market Plaza, Steuart St. Tower,
    STREET: Twentieth Floor
    CITY: San Francisco
              CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND AND TOWNSEND KHOURIE and CREW
STREET: One Market Plaza, Steuart St. Tower,
STREET: Twentieth Floor
CITY: San Francisco
                                                                                                                                                  COUNTRY: USA
ZIP: 94105-1492
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/453,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                          PPLICATION CATALLY PILLING DATE:

CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/100,887

FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: PAIMER: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: PAIMER: 31-990
REGISTRATION NUMBER: 31-990
REGISTRATION NUMBER: 31-990
REFERENCE/DOCKET NUMBER: 13952-15-1
TELEPHONE: 206-467-9600
; TELEPHONE: 206-467-9600
; TELEPRATION INFORMATION:
; TELEPRATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
remaint 44 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8;
Pred. No. 43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 GAATTCACTAGTGAGCTCTT 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 GAACUCACUCGUGAGCUCTT 22
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70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 70.0
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 44 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 94105-1492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: line
| IMMEDIATE SOURCE
| CLONE: ZC2938
US-08-453-742-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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12 GAATTCACTAGTGAGCTCTT 31
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                                                                                                                                                        Score 16.8; DB 1; Length 44; Pred. No. 43;
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Pred. No. 43;
4; Mismatches 2; Indels
                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                     Sequence 15, Application US/08452802

Patent No. 5683884

GENERAL INFORMATION:
APPLICANT: Moore, Emma E
APPLICANT: Kuestner, Paul O
APPLICANT: Kuestner, Rolf E
ITILE OF INVENTION: Human Calcitonin Receptor
NUMBER OF SEQUENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND AND TOWNSEND KHOURIE and CREW
STREET: One Market Plaza, Steuart St. Tower,
CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 30-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/954,804
FILING DATE: 02-AUG-1993
APPLICATION NUMBER: US 07/954,804
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: PARTHELS, Steven W
REGISTATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 13952-15-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                     3 GAACUCACUCGUGAGCUCTT 22
                                                                                                                                                                                                                                        12 GAATTCACTAGTGAGCTCTT 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.4%;
                                                                                                                                                        Query Match
Best Local Similarity 70.0%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : 206-467-9600
415-543-5043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 70.0°
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-543-5043
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
                                              TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                        IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IMMEDIATE SOURCE:
                                                                                                        US-08-453-222-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                             US-08-452-802-15
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3 GAACUCACUCGUGAGCUCTT 22

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Sequence 38, Application US/08479939
; Father No. 5686593
; Pather No. 5686593
; GENERAL INFORMATION:
APPLICANT: Woeldike, Helle F.
APPLICANT: Woeldike, Helle F.
APPLICANT: Hjort, Carsten M.
APPLICANT: Sven, Hagen, Frederick
APPLICANT: Sven, Hagen, Engyme Capable of Degrading Cellulose
; TITLE OF INVENTION: or Hemicellulose
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 44;
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                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: United States of America
ZIP: 10174-6201
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/479,939
FILING DATE: 07-UN-1995
CLASSIFICATION NUMBER: US/08/361,920
FILING DATE: 22-DEC-1994
APPLICATION NUMBER: US 07/940,860
FILING DATE: 28-OCT-1992
APPLICATION NUMBER: US 07/940,860
FILING DATE: 08-MAY-1990
RION APPLICATION DATA:
APPLICATION NUMBER: DK 1158/90
FILING DATE: 08-MAY-1991
ATTORNEY/AGENT INFORMATION:
NUMBER: LAMBLICATION NUMBER: PCT/DK91/00124
FILING DATE: 08-MAY-1991
ATTORNEY/AGENT INFORMATION:
NUMBER: LAMBLICATION NUMBER: DK 1158/100124
FILING DATE: 08-MAY-1991
ATTORNEY/AGENT INFORMATION:
NUMBE: LAMBLICATION NUMBER: DK 1158/100124
FILING DATE: 08-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ed. No. 43;
Mismatches
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Conservative 4; Mismatche
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APPLICANT: Woeldike, Helle F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 33,728
REPERBNEK/GOCKET UNBER: 34.
TELECOMMUNICATION:
TELEPHONE: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: DNA (genomic)
US-08-479-939-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
---- 14; Conserve
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                                                                                                                                                                                                                                                                                                                                                           CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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12 GAATTCACTAGTGAGCTCTT 31
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                                                                                                                                                              SEE: No. 57632540 No. 5763254disk of No. 5763254th America, Inc. : 405 Lexington Avenue, 62nd Floor New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
APPLICANT: Hjort, Carsten M.
APPLICANT: Sven, Hastrup
TITLE OF INVENTION: or Hemicellulose
NUMBER OF SEQUENCES: 85
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
ADDRESSEE: NO. 57632540 NO. 5763254th Am
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 70.0%; Pred. No. 43;
Matches 14; Conservative 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC Compatible
OPERATING SYSTEM:
PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,432
FLING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 07/940,860
FLING DATE: 28-OCT-1992
APPLICATION NUMBER: US 07/940,860
FLING DATE: 28-OCT-1992
APPLICATION NUMBER: DY 1158/90
FILING DATE: 09-MAY-1990
PRIOR APPLICATION DAMBER: PCT/DK91/00124
FILING DATE: 08-MAY-1991
APPLICATION NUMBER: PCT/DK91/00124
FILING DATE: 08-MAY-1991
APPLICATION NUMBER: 33.728
REGISTRATION NUMBER: 33.728
REGISTRATION NUMBER: 3435.204-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lehner, Joyce M.
TITLE OF INVENTION: MAMMALIAN ZCYTORS
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Zymogenetics
                                                                                                                                                                                                                                        STATE: New York
COUNTRY: United States of America
ZIP: 10174-6201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12, Application US/09071224;
Patent No. 6271343
GENERAL INFORMATION:
APPLICANT: Lok, Si
APPLICANT: Presnell, Scott R.
APPLICANT: Jelmberg, Anna C.
APPLICANT: Gilbert, Teresa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 GAATTCACTAGTGAGCTCTT 31
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Presnell, Scott R.
Jelmberg, Anna C.
Gilbert, Teresa
Foster, Donald C.
Adams, Robyn L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zymogenetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-483-432-38
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 15
US-09-071-224-12
                                                                                                                                                                                         STREET:
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Gaps
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76.4%; Score 16.8; DB 3; Length 44;
Best Local Similarity 70.0%; Pred. No. 43;
Matches 14; Conservative 4; Mismatches 2; Indels
                                                                                                           COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
SOFTWARE: FASTESO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,224
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
FILING DATE:
ATMORMEY/AGENT INPORMATION:
NAME: LUND, Paul G
                    1201 Eastlake Ave East
                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Lunn, Paul G
REGISTRATION NUMBER: 32,743
REPRENCE/DOCKET NUMBER: 96-2:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6627
TELEFAX: 206-442-6678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 GAACUCACUCGUGAGCUCTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
STREET: 1200
                                                                                  USA
                                                                                                    98102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-071-224-12
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Search completed: June 24, 2005, 04:11:08 Job time : 64.4724 secs

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90.9 24774
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                                                                                                                                       June 24, 2005, 00:03:32 ; Search time 253.433 Seconds (without alignments) 542.594 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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/cgn2_6/ptodata1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata1/pubpna/US06_NEW_PUB.seq:*
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/cgn2_6/ptodata1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata1/pubpna/US096_PUBCOMB.seq:*
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                 GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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                                                                                               - nucleic search, using sw model
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Maximum DB seq length: 200000000
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22
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		Description		Sequence 1, Appli	Seguence 780, App	Sequence 29766, A	Sequence 4, Appli	Sequence 14, Appl	Sequence 3, Appli	Sequence 29748, A
SUMMARIES		. QI		US-10-848-737-1	US-10-831-901A-780	US-10-831-901A-29766	US-10-699-936-4	US-10-699-936-14	US-10-889-447-3	US-10-831-901A-29748
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B 20 90.9 24774 21 10 20 90.9 28920 21 11 20 90.9 28920 21 12 20 90.9 28920 21 13 20 90.9 28920 21 14 20 90.9 28920 21 15 20 90.9 28920 21 16 20 90.9 28920 21 17 20 90.9 29291 21 18 20 90.9 29291 21 18 20 90.9 29291 21 20 20 90.9 29291 21 21 20 90.9 29291 21 22 20 90.9 29291 21 23 20 90.9 29705 21 24 20 90.9 29705 21 25 20 90.9 29705 21 26 20 90.9 29705 21 27 20 90.9 29711 21 28 20 90.9 29711 21 28 20 90.9 29712 21 39 20 90.9 29712 21 31 20 90.9 29711 21 32 20 90.9 29712 21 33 20 90.9 29712 21 34 20 90.9 29712 21 35 20 90.9 29712 21 36 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712 21 39 20 90.9 29712 21 39 20 90.9 29712 21 31 20 90.9 29712 21 32 20 90.9 29712 21 34 20 90.9 29712 21 35 20 90.9 29712 21 36 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712 21 39 20 90.9 29712 21 31 20 90.9 29712 21 32 20 90.9 29712 21 34 20 90.9 29712 21 35 20 90.9 29712 21 36 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712 21 38 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712 21 38 20 90.9 29712 21 39 20 20 20 20 20 30 29712 21 31 20 90.9 29712 21 32 20 90.9 29712 21 34 20 90.9 29712 21 37 20 90.9 29712 21 38 20 90.9 29712	889-101-3 889-447-5 881-9017-5 889-101-6 889-101-6 889-101-7 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29 881-9017-29	ALIGNMENTS 37 5 SARS-ASSOCIATED ND REPLICATION BY 1/848,737 1/901
B 20 90.9 24774 10 20 90.9 28920 11 20 90.9 28920 11 20 90.9 28920 12 20 90.9 28920 13 20 90.9 28920 14 20 90.9 28920 15 20 90.9 28920 16 20 90.9 29033 16 20 90.9 29291 18 20 90.9 29291 20 20 90.9 29291 21 20 90.9 29430 22 20 90.9 29430 23 20 90.9 29430 24 20 90.9 29705 25 20 90.9 29705 26 20 90.9 29705 27 20 90.9 29705 28 20 90.9 29705 28 20 90.9 29705 29 20 90.9 29705 29 20 90.9 29705 29 20 90.9 29705 2		11. (1. (1. (1. (1. (1. (1. (1. (1. (1.
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B 8 20 9 9 20 10 20 11 20 11 20 12 20 13 20 14 20 15 20 16 20 17 20 18 20 19 20 20 20 21 20 21 20 22 20 23 20 24 20 26 20 27 20 28 20 29 20 31 20 31 20 32 20 33 20 34 20 36 20 37 20 38 20 39 20 30 20 31 20 31 20 32 20 34 20 36 20 37 20 38 20 39 20 30 20 30 20 31 20 31 20 32 20 33 20 34 20 36 20 37 20 38 20 39 20 30 20 30 20 31 20 30 20 31 20 32 20 33 20 34 20 36 20 37 20 38 20 39 20 30 20 30 20 31 20 32 20 33 20 34 20 36 20 37 20 38 20 39 20 30 20 30 20 30 20 30 20 30 20 30 20 30 20 30 20 45 20 45 20 46 20 47 20 47 20 48 20 49 20 40 20	000000000000000000000000000000000000000	plicat:
RESULT 1 11.1 11.1 11.1 11.1 11.1 11.1 11.1 1		137-1 1, Ap 11, Ap 11, Ap 11: Ba 11:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sampath, Rangarajan
APPLICANT: Sampath, Rangarajan
APPLICANT: Freier, Susan M.
APPLICANT: Freier, Susan M.
APPLICANT: Hofstadler, Steven A.
APPLICANT: Hofstadler, Steven A.
APPLICANT: Hofstadler, Steven A.
APPLICANT: Lowery, Kristin Sannes
APPLICANT: Savaye, Eric
APPLICANT: Savaye, Eric
APPLICANT: Baker, Brenda F.
APPLICANT: Bennett, C. Frank
TITLE OF INVENTION: Compositions And Methods For The Treatment Of Severe
TITLE OF INVENTION: Acute Respiratory Syndrome (SARS)
FILE REFERENCE: ISISO083-100 (BIOL0008US)
CURRENT APPLICATION NUMBER: US/10/831,901A
CURRENT APPLICATION NUMBER: 60/466,426
PRIOR FILING DATE: 2003-04-28
PRIOR FILING DATE: 2003-06-66
PRIOR PRILING DATE: 2003-06-66
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR PRILING DATE: 2003-06-27
NUMBER OF SEQ ID NOS: 30063
SCOTT NO NO NOS: 30063
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                                           Query Match
100.0%; Score 22; DB 21; Length 22;
Best Local Similarity 100.0%; Pred. No. 0.37;
Matches 22; Conservative 0; Mismatches 0; Indels
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Publication No. US20050100885A1
GENERAL INFORMATION:
APPLICANT: Crooke, Stanley T.
APPLICANT: Ecker, David J.
APPLICANT: Fraier, Sumpath, Rangarajan
APPLICANT: Fraier, Susan
APPLICANT: Hofstadler, Steven A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ), OTHER INFORMATION: Antisense compound US-10-831-901A-780
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; Sequence 780, Application US/10831901A
; Publication No. US20050100885A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Crooke, Stanley T. APPLICANT: Ecker, David J.
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Matches 15; Conserv
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LENGTH: 20
US-10-848-737-1
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Squence 4, Application US/1069936

Squence 4, Application US/1069936

Squence 4, Application US/1069936

Squence 4, Application Wo. US205009582A1

GENERAL INFORMATION:
APPLICANT: Gilliam Ross, Laura
APPLICANT: Scholl, David R.
APPLICANT: Wentworth, David R.
APPLICANT: Wentworth, David E.
APPLICANT: O'Ollick, Joseph D.
TITLE OF INVENTION: Compositions and Methods for Detecting Severe Acute Respiratory
TITLE OF INVENTION: Syndrome Coronavirus
FILE REPERENCE: DHI-O'7986

CURRENT FILING DATE: 2003-11-03

NUMBER OF SEQ ID NOS: 87

SOFTWARE: Patentin version 3.2

SEQ ID NO 4

LENGTH: 1706

MANDER OF THE TOR OF THE 
APPLICANT: Swayze, Eric
APPLICANT: Baker, Brenda F.
APPLICANT: Baker, Brenda F.
APPLICANT: Bannett, C. Frant
TITLE OF UNUENTION: Compositions And Methods For The Treatment Of Severe
TITLE OF INVENTION: Acute Respiratory Syndrome (SARS)
FILE REFERENCE: ISTSO033-100 (BIOLO008US)
CURRENT FILING DATE: 2004-04-26
PRIOR APPLICATION NUMBER: 60/466,426
PRIOR FILING DATE: 2003-04-28
PRIOR PILING DATE: 2003-05-06
PRIOR APPLICATION NUMBER: 60/468,562
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-10
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PRIOR PILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-27
PRIOR PILING DATE: 2003-06-27
NUMBER OF SEO ID NOS: 3006-3
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 29766
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Pred. No. 4.3;
5; Mismatches 0; Indels
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US-10-699-936-14
; Sequence 14, Application US/10699936
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Best Local Similarity 75.0%;
Matches 15; Conservative
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Best Local Similarity 75.0%;
Matches 15; Conservative
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NAME/KEY: misc_feature

| NAME/KEY: misc_feature

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| LOCATION: 4134-4233, 4562-4662, 5012-5111, 5781-5879, 7438-7537, 7837-7936,

| LOCATION: 816-8715, 12025-12124, 13984-14083, 16463-16562, 16932-17031,

| LOCATION: 17381-17480, 18090-18189, 19019-19118, 19478-19577, 20357-20456,

| LOCATION: 21086-21185, 21945-22044, 23174-23273, 23531

| US-10-831-901A-29748
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Pred. No. 4.6;
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                                                                           APPLICANT: SWAYZE, EXIC
APPLICANT: BAREY, BEENER
APPLICANT: BAREY, BEENER
APPLICANT: BAREY, BEENER
TITLE OF INVENTION: Compositions And Methods For The Treat
TITLE OF INVENTION: Compositions And Methods For The Treat
TITLE OF INVENTION: Composition of BIOLOGOGUS)
FILE REFERENCE: ISISO083-100 (BIOLOGOGUS)
CURRENT APPLICATION NUMBER: US/10/831,901A
CURRENT PILING DATE: 2004-04-26
PRIOR FILING DATE: 2003-04-26
PRIOR FILING DATE: 2003-06-06
PRIOR FILING DATE: 2003-06-06
PRIOR FILING DATE: 2003-06-06
PRIOR FILING DATE: 2003-06-06
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-27
PRIOR FILING DATE: 2003-06-27
PRIOR FILING DATE: 2003-06-27
NUMBER OF SEQ ID NOS: 30063
SEQ ID NO 29748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/10889101
Fublication No. US20050107324A1
GENERAL INFORMATION:
APPLICANT: Bennett, C. Frank
APPLICANT: Dobie, Kenneth W.
TITLE OF INFORMATION: MODULATION OF CEACAM1
FILE REFERENCE: ISISO101-100 (RTS-0655US)
CURRENT APPLICATION NUMBER: US/10/889,101
CURRENT FILING DATE: 2004-07-12
FRIOR APPLICATION NUMBER: 60/486,652
FRIOR APPLICATION NUMBER: 60/486,652
FRIOR FILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 298
SOFTWARE: FREESEQ FOR WINDOWS VERSION 4.0
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706 GTGAACTCACTCGTGAGCTC 725
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Best Local Similarity 75.0%;
Matches 15; Conservative
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LENGTH: 24774
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                                                            APPLICANT: Taylor, Jill
APPLICANT: Scholi, David R.
APPLICANT: Scholi, David R.
APPLICANT: Wentworth, David E.
APPLICANT: Wentworth, David E.
APPLICANT: Wentworth, David E.
TITLE OF INVENTION: Compositions and Methods for Detecting Severe Acute Respiratory;
FILE REPRENCE: DHI-07986
CURRENT APPLICATION NUMBER: US/10/699,936
CURRENT FILING DATE: 2003-11-03
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PatentIn version 3.2
SEQ ID NO 14
LENGTH: 1706
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; Publication No. US20050075307A1
; GENERAL INFORMATION:
   APPLICANT: Beneric C. Frank
   APPLICANT: Jain, Ravi
   TILE OF INVENTION: MODULATION OF AMINOPEPTIDASE N EXPRESSION
   TILE OF INVENTION: MODULATION OF AMINOPEPTIDASE N EXPRESSION
   CURRENT ENERGES: 2004-07-12
   CURRENT FILING DATE: 2004-07-12
   PRIOR APPLICATION NUMBER: 60/486,670
   PRIOR FILING DATE: 2003-07-12
   NUMBER OF SEQ ID NOS: 241
   SOFTWARE: FastSEQ for Windows Version 4.0
   SEQ ID NO 3
   LENGTH: 24774
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; ORGANISM: SARS coronavirus Shanghai LY
US-10-699-936-14
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; OTHER INFORMATION: n is any nucleotide
US-10-889-447-3
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APPLICANT: Ecker, David J.
APPLICANT: Sampath, Rangarajan
APPLICANT: Freier, Susan M.
APPLICANT: Massire, Christian
US20050095582A1
                                                      APPLICANT: Gillim-Ross, Laura APPLICANT: Taylor, Jill
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Best Local Similarity 75.0
Matches 15; Conservative
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APPLICANT: Bennett, C. Frank
TITLE OF INVENTION: Compositions And Methods For The Treatment Of Severe
TITLE OF INVENTION: Acute Respiratory Syndrome (SARS)
FILE REPERENCE: ISISOOBBA3-100 (BIOL0008US)
CURRENT APPLICATION NUMBER: US/10/831,901A
CURRENT FILING DATE: 2004-04-26
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) FRATURE:

NAME/KEY: misc feature

| LOCATION: 7230-7329, 9929-10028, 10137-10236, 11646-11745, 20385-20484,

| LOCATION: 21024-21123, 21753-21852, 22112-22211, 25301-25400

| COTHER INFORMATION: n = A,T,C or G

US-10-831-991A-29740
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; Publication No US20050107324A1
; GENERAL INFORMATION:
APPLICANT: Bennett, C. Frank
APPLICANT: John, Ravi
TILLE OF INVENTION: MODULATION OF CEACAM1 EXPRESSION
FILE REFERENCE: ISISO101-100 (RTS-0655US)
CURRENT APPLICATION NUMBER: US/10/889,101
; CURRENT PILING DATE: 2004-07-12
; PRIOR FILING DATE: 2003-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/466,426
PRIOR PILING DATE: 2003-04-28
PRIOR PILING DATE: 2003-04-28
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-04-30
PRIOR PILING DATE: 2003-04-30
PRIOR PILING DATE: 2003-06-06
PRIOR PILING DATE: 2003-06-06
PRIOR PILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-17
NUMBER OF SEQ ID NOS: 30063
SOFTWARE: PRESEES FOR WINDOWS VERSION 4.0
SEQ ID NO 29740
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                                                                                                                                       Sequence 29740, Application US/10811901A Publication No. US20050100885A1 GENERAL INFORMATION:
APPLICANT: Crooke, Stanley T. APPLICANT: Ecker, David J.
  776 GTGAACTCACTCGTGAGCTC 795
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776 GTGAACTCACTCGTGAGCTC 795
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Lowery, Kristin Sannes
Swayze, Eric
Baker, Brenda F.
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Freier, Susan M.
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Pred. No. 4.6;
  DB 21; Length 24774;
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TITLE OF INVENTION: MODULATION OF AMINOPEPTIDASE N EXPRESSION
FILE REFERENCE: RTS-0665US
CURRENT APPLICATION NUMBER: US/10/889,447
CURRENT PILING DATE: 2004-07-12
PRIOR APPLICATION NUMBER: 60/486,670
PRIOR PILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 241
SOFTWARE: FRRESEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 28920
                                                                                                                                                                                                                                                                                                                              APPLICANT: Bennett, C. Frank
APPLICANT: Jain, Ravi
TITLE OF INVENTION: MODULATION OF AMINOPEPTIDASE N EXPRESSION
TITLE OF INVENTION: MODULATION OF AMINOPEPTIDASE N EXPRESSION
CURRENT APPLICATION NUMBER: US/10/889,447
CURRENT FILING DATE: 2004-07-12
FRIOR APPLICATION NUMBER: 60/486,670
FRIOR APPLICATION NUMBER: 60/486,670
NUMBER OF SEQ ID NOS: 241
SOFTWARE: FastSEQ for Windows Version 4.0
                                                 0; Indels
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5; Mismatches
Query Match 90.9%; Score 20; DB Best Local Similarity 75.0%; Pred. No. 4.6; Matches 15; Conservative 5; Mismatches
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ORGANISM: SARS coronavirus isolate BJ03
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                                                                                                                        706 GTGAACTCACTCGTGAGCTC 725
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776 GTGAACTCACTCGTGAGCTC 795
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; Sequence S, Application US/10889447
; Publication No. US20050075307A1
; GENERAL INFORMATION:
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Best Local Similarity 75.0%;
Matches 15; Conservative !
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Best Local Similarity 75.uv
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LENGTH: 28920
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NUMBER OF SEQ ID NOS: 298

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CURRENT APPLICATION NUMBER: US/10/831,901A
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Best Local Similarity
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                                                                                                                                                                                                                                        DB 21; Length 28920;
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APPLICANT: Dobie, Kenneth W.
APPLICANT: Jain, Ravi
APPLICANT: Jain, Ravi
TITLE OF INVENTION: MODULATION OF CEACAMI EXPRESSION
FILE REFERENCE: ISISO101-100 (RTS-0655US)
CURRENT APPLICATION NUMBER: US/10/889,101
CURRENT FILING DATE: 2004-07-12
PRIOR APPLICATION NUMBER: 60/486,652
PRIOR APPLICATION NUMBER: 60/486,652
PRIOR FILING DATE: 203-07-12
NUMBER OF SEQ ID NOS: 298
SOFTWARE: FRSESQ IO NOS: 298
                                                                                                                                                                                                                                                                                5; Mismatches
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                      Score 20;
Pred. No.
                                                                                    ORGANISM: SARS Coronavirus isolate BJ03
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ORGANISM: SARS Coronavirus isolate BJ04
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APPLICANT: Crooke, Stanley T. APPLICANT: Ecker, David J.
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                                                                                                                    NAME/KEY: misc_feature; LOCATION: COTHER INFORMATION: n = A,T,C or GUS-10-889-101-5
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US-10-889-101-6
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Hofstadler, Steven A.
Lowery, Kristin Sannes
Swayze, Eric
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Freier, Susan M.
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Best Local Similarity 75.0%;
Matches 15; Conservative
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                    SEQ ID NO 5
LENGTH: 28920
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LENGTH: 28920
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                                                                TYPE: DNA
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APPLICANT: Massire, Christian
APPLICANT: Hofstadler, Steven A.
APPLICANT: Hofstadler, Steven A.
APPLICANT: Lowery, Kristin Sannes
APPLICANT: Swayze, Eric
APPLICANT: Baker, Brenda F.
APPLICANT: Bennett, C. Frank
TITLE OF INVENTION: Compositions And Methods For The Treatment Of Severe
TITLE OF INVENTION: Acute Respiratory Syndrome (SARS)
FILE REFERENCE: ISISO003-100 (BIOL0008US)
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Pred. No. 4.6;
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CURRENT FILING DATE: 2004-04-26
PRIOR PRICATION NUMBER: 60/466,426
PRIOR PILING DATE: 2003-04-28
PRIOR PILING DATE: 2003-05-06
PRIOR PILING DATE: 2003-05-06
PRIOR PELING DATE: 2003-05-06
PRIOR PELING DATE: 2003-04-30
PRIOR PELING DATE: 2003-04-30
PRIOR PELING DATE: 2003-06-06
PRIOR PILING DATE: 2003-06-06
PRIOR PILING DATE: 2003-06-06
PRIOR APPLICATION NUMBER: 60/477,637
PRIOR APPLICATION NUMBER: 60/477,637
PRIOR APPLICATION NUMBER: 60/477,637
PRIOR APPLICATION NUMBER: 60/483,579
PRIOR PILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: 60/466,426
PRIOR FILING DATE: 2003-04-29
PRIOR FILING DATE: 2003-04-29
PRIOR FILING DATE: 2003-05-06
PRIOR FILING DATE: 2003-05-06
PRIOR FILING DATE: 2003-05-06
PRIOR FILING DATE: 2003-06-10
PRIOR PILING DATE: 2003-06-10
PRIOR APPLICATION NUMBER: 60/469,627
PRIOR APPLICATION NUMBER: 60/483,579
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
PRIOR SELING DATE: 2003-06-10
PRIOR FILING DATE: 2003-06-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 29742, Application US/10831901A Publication No. US20050100885A1
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APPLICANT: Ecker, David J.
APPLICANT: Sampath, Rangarajan
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Best Local Similarity 75.0%;
Matches 15; Conservative
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US-10-831-901A-29819
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ORGANISM: SARS Coronavirus
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0; Gaps

Search completed: June 24, 2005, 04:35:43 Job time: 259.433 secs

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June 23, 2005, 23:49:32; Search time 1551.26 Seconds (without alignments) 539.828 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                68479088
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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9b htc::22

9b est4::3

9b est4:3:3

9b 9s8:3

9b 9s8:1:3
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Perfect score:
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•	Description	AQ479024 RPCI-11-2	BG917388 602816903	BF100955 601754406		AZ312688 1M0028J02	BP091999 BP091999	BQ820183 1030082F1	AZ236076 RPCI-23-8	AV389206 AV389206	CB213780 OML04060	AV619265 AV619265	AZ259445 RPCI-23-1	BQ824823 1030121E1	AZ695771 RPCI-23-2	BQ820182 1030082F1	BG844655 1024007C0	AG427536 Mus muscu	BF862564 963036A08	AQ997054 RPCI-23-3	AQ996162 RPCI-23-3	CN680938 E0144A04-	CE718342 tigr-gss-	CE779822 tigr-gss-	BF163602 601769987
SUMMARIES	ID	AQ479024	BG917388	BF100955	AK031386	AZ312688	BP091999	BQ820183	AZ236076	AV389206	CB213780	AV619265	AZ259445	BQ824823	AZ695771	BQ820182	BG844655	AG427536	BF862564	AQ997054	AQ996162	CN680938	CE718342	CE779822	BF163602
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de	Query Match	85.5	85.5	83.6	83.6	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80.9	80:9	80.9	79.1	79.1	79.1	79.1	79.1	79.1
	Score	18.8	18.8	18.4	18.4	17,8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.8	17.4	17.4	17.4	17.4	17.4	17.4
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/cell_type="Lymphocytes" /clone lib="RPC1-11" /noce="Vector: pBACe3.6; Site_1: BcoR1; Site_2: BcoR1; RPCI11 Human Male BAC Library"

ORIGIN

.4 79.1 712 5 BU709067 BU709067 .4 79.1 732 9 AG401233 AG401233 .4 79.1 919 9 CL501597 CL501597 .2 78.2 35 7 ZM466607 AM466600 .2 78.2 355 7 CR473310 CR475310 .2 78.2 385 1 AA087904 AA087904 .2 78.2 446 9 CL3569486 CL359486 .2 78.2 458 1 AL924913 AL924913	17.2 78.2 46.2 4 B6398975 B63388975 B6398975 B6398977 B639978 B639978	ALIGNMENTS	AQ479024 RPCI-11- RPCI-11- RPCI-11- RQ479024 AQ479024 AQ4	/www_rype=_genomiry /db_xref="qfb97444" /db_xref="taxon:9606" /clone="RRCI-11-254MS" /cell type="Lymphocytes"
C C 239 330 330	00 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0		RESULT 1 AQ479024 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE OUGGANISM REFERENCE AUTHORS TITLE JOURNAL COMMENT COMMENT FEATURES BOUICE	

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                                                                                                                                                                                                                         BG917388 1344 bp mRNA linear EST 05-JUN-2001
602816903F1 NCI_CGAP_Mam6 Mus musculus cDNA clone IMAGE:4945921 5',
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/sex="female, virgin"
/tissue trype="infiltrating ductal carcinoma"
/tissue trype="infiltrating ductal carcinoma"
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/clone_lib="NCI_GGAP_Mam6"
/note="Organ: mammarry; Vector: pCMV-SPORT6; Site 1: Sal1;
Site_2: Not1; Cloned unidirectionally. Primer: Öligo dT.
Library constructed by Life Technologies. Investigator
providing samples: Jeffrey Green, M.D., NIH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                         National institutes of health, Mammalian Gene Collection (MSL)
Uppublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Jeffrey Green M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LiNL at:
http://image.llh.gov
Plate: LiLAM10893 row: f column: 02
High quality sequence stop: 309.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                      Mus musculus
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (Dases 1 to 1344)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                     Length 254;
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                   Score 18.8; DB 8;
Pred. No. 1.3e+02;
5; Mismatches 2;
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996 GTGGTCTCACTCGTGAGCTCTT 975
                                                                                            22
                                                                                                               1 GUGAACUCACUCGUGAGCUCTT 22
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/strain="FVB/N"
                                                                                            1 GUGAACUCACUCGUGAGCUCTT
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BG917388.1 GI:14297864
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                   85.5%;
68.2%;
Query Match
Best Local Similarity 68.2v
Best Local Similarity 68.2v
-hos 15; Conservative
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/strain="kype="mRNA"
/strain="kype]Nu"
/db_xref="taxon:10090"
/clone="IMAGE:3983736"
/tissue type="tumor, biopsy sample"
/dev stage="3 months, virgin"
/dev stage="3 months, virgin"
/dev stage="3 months, virgin"
/dev stage="1 months, v
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llni.gov
Plate: LLAM9184 row: k column: 01

High quality sequence stop: 579.
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Musinae, Mus
1 (bases 1 to 915)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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High-efficiency full-length cDNA cloning
Weth. Enzymol. 303, 19-44 (1999)
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/note="unnamed protein product; DJ1014D13.2 (NOVEL PROTEIN
SIMILAR TO ACTN3 (ACTININ, ALPHA 3)) (FRAGMENT) homolog
[Homo sapiens] (SPTR|Q9UH44, evidence: FASTY, 88.7%ID,
65.2%length, match=372)
Shibata, K., Itoh, M., Aizawa, K.; Nagaoka, S., Sasaki, N., Carninci, P., Konno, H., Akiyama, J., Nishi, K., Kitsunai, T., Tashiro, H., Itoh, M., Sumi, N., Ishii, Y., Nakamura, S., Hazama, M., Nishine, T., Harada, A., Yamamoto, R., Matsumoto, H., Sakayuchi, S., Ikegami, T., Kashiwagi, K., Fujiwake, S., Inoue, K., Togawa, Y., Izawa, M., Ohara, B., Watahiki, M., Yoneda, Y., Ishikawa, T., Ozawa, K., Tanaka, T., Matsuura, S., Kawai, J., Okazaki, Y., Muramatsu, M., Inoue, Y., Kira, A. and Hayashizaki, Y. RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (16-UL-2001) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/, Tel:81-45-503-9222,
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/clone_lib="RIXEN full-length enriched mouse cDNA library"
/dev_stage="13 days embryo"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The FANTOM Consortium and the RIKEN Genome Exploration Research Group Phase I & II Team.
Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs
Nature 420, 563-573 (2002)
6 (bases 1 to 1580)
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                                                                                                                                                                                                                                                                                                                                                                                                      The RIKEN Genome Exploration Research Group Phase II Team and the
                                                                                                                                                                                                                                                                                                                                                                                                                                  FANTOM Consortium. Functional annotation of a full-length mouse cDNA collection Nature 409, 685-690 (2001)
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Please visit our web site for further details.
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'db_xref="taxon:10090"
clone="6030419L17"
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/protein_id="BAC27378.1"
/db_xref="G1:26327269"
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URL:http://fantom.gsc.riken.jp/.
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/strain="C57BL/6J"
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/lab host="E. Coli strain Xi10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42IV; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resources/documents/dhares/): The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end repaired with T4 DNA polymerses and T4
polymuclectide kinase. Adaptor oligonuclectides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
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/translation="magprgallawcrrqcegyrgydirdlsssfrdglafcailhrh
rpdlldfqslskenvfennrlafevarkelgipalldpndmvsmsvpdclsimtyvsq
yynhftssgqgergpspsgrldmavyffaeq"
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1M0028J02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0028J02 R, genomic survey sequence.
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                                                                                                                                                                                                                                                                           Length 1580;
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Pred. No. 2.8e+02;
5; Mismatches 1;
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Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah Genome Center
University of Utah
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/strain="C57BL/6J"
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/clone="UUGC1M0028J02"
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
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776 GTGAACTCACTGGTGAGCTC 757
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Matches 14; Conservative
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Phycologia (2004) In press
Contact: Erika Asamizu
The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute
Yana 153-3. Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.ip, URL:http://www.kazusa.or.jp/en/plant/.
Location/Qualifiers
adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chlamydomonadaceae; Chlamydomonas.

(bases 1 to 506)

Asanizu, E., Nakamura, Y., Miura, K., Fukuzawa, H., Fujiwara, S.,
Hirono, M., Iwamoto, K., Matsuda, Y., Minagawa, J., Shimogawara, K.,
Takahashi, Y. and Tabata, S.
Establishment of Publicly Available cDNA Material and Information
Resource of Chlamydomonas reinhardtii (Chlorophyta), to Facilitate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BQ820183 514 bp mRNA linear EST 01-AUG-2005
1030082F11.y2 C. reinhardtii CC-1690, Deflagellation (normalized),
Lambda Zap II Chlamydomonas reinhardtii cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                     BP091999 Chlamydomonas reinhardtii C9 various conditions
Chlamydomonas reinhardtii cDNA clone MX248b10_r 5', mRNA sequence.
BP091999
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Chlamydomonas reinhardtii
Eukaryota; Viridiplantae; Chlorophyta; Chlorophyceae; Volvocales;
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                                                                                             Length 451;
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/mol_type="mRNA"
/strain="C9"
                                                                                         Score 17.8; DB 8;
Pred. No. 4.7e+02;
4; Mismatches 2;
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                                                                                             80.9%;
71.4%;
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Best Local Similarity 71.4
Matches 15, Conservative
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Best Local Similarity 66.7
Matches 14; Conservative
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/Mol type="mrkNa"
//wol type="mrkNa"
//wol type="mrkNa"
//wol type="mrkNa"
//wormalized.1650 wild type mt + 21gr"
//wormalized.build type mt + 21gr"
//wormalized), lambda Zap II"
//noce="wetor: pBluescript II SK-; Site_!: EcoR!; Site_2:
//noce="wetor: pBluescript II SK-; Site_!: EcoR!; Site_2:
//noce="wetor: pBluescript II SK-; Site_1: EcoR!; Site_2:
//which had been re-synthesizing flagellated by John Davies
//which had been re-synthesizing flagellated by John Bond 60
//which had been re-synthesizing flagellated by PH shock. PolyA mRNA was
purified from each sample, pooled and cDNA synthesized
//wormally cloned into lambda Zap II
//stratagene) in the EcoR! (5') and Xhor! (3') sites.
//pBluescript II SK- plasmids were excised from the lambda
//wormalized using method 4 described
in Bonaldo et al., (1996) Genome Research 6: 791-806."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eu (bases 1 to 514)

S. 1 (bases 1 to 514)

S. Ano,S., Nierman,W., Feldblyum,T., Malek,J., Shatsman,S.,
Akinret,B., Levins,M., Mcgann,S., Tsegaye,G., Geer,K., Krol,M., de
Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-23

Unpublished (1999)
Other GSSs: RPCI-23-84D17.TV
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, USA
Tel: 301 838 0208
E 1 (bases 1 to 514)

S Grossman, A., Chang, C.-W., Davies, J., Harris, E., Hauser, C.,
Lefebyre, P., McDernott, J.P., Shrager, J., Silfow, C. and Stern, D.
Lefebyre, P., McDernott, J.P., Shrager, J., Silfow, C. and Stern, D.
Analyses of the Chlamydoonas reinhardtii Genome: A Model,
Unicellular System for Analyzing Gene Function and Regulation in
Vascular Plants. Project: 1030

L Unpublished (2002)

Contact: Charles Hauser

DCMB Box 91000

Duke University

Durham, NC 27708-1000

Tel: 919 613 8159

Fax: 919 613 8177

Email: chauser@duke.edu.

Location/Qualifiers
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RPCI-23-84D17.TJ RPCI-23 Mus musculus genomic clone RPCI-23-84D17, 99nomic survey sequence.
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Clones are derived from the mouse BAC library RPCI-23. For BAC
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/clone lib="RPCI-23"
/clone lib="RpCI"
/clone li
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AV389206 AV389206 GI:6543422
EST.
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library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/orderingframe.htm)
or from Resea ch Genetics (info@resgen.com). BAC end page:
http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Plate: 84 row: D column: 17
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The First Laboratory for Plant Gene Research
Azusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: ynakamu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/
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Asamizu, E., Nakamura, Y., Sato, S., Pukuzawa, H. and Tabata, S.
A large scale structural analysis of cDNAs in a unicellular green alga, Chlamydomonas reinhardtii. I. Generation of 3433
DNA Res. 6 (6), 369-373 (1999)
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Chlamydomonadaceae, Chlamydomonas.
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/clone="CM040a08 r"
/dev stage="photoautotrophic growth"
/clone_lib="Chlamydomonas reinhardtii C9"
/note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2:
Xho!="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2:
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    544
/organism="Chlamydomonas reinhardtii"

                                                                                                                                                                                                                                                                                                                 'organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                       /mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="RPCI-23-84D17"
                                                                                                                                                                                                                 Class: BAC ends.
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/strain="C9"
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Chlamydomonas reinhardtii
Chlamydomonas reinhardtii
Chlamydomonas reinhardtii
Chlamydomonas reinhardtii
Chlamydomonas.

1 (bases 1 to 546)
Asamizu, E., Mutra, K., Kucho, K., Inoue, Y., Fukuzawa, H., Ohyama, K., Nakamura, Y. and Tabata, S.
Generation of expressed sequence tags from low-CO2 and high-CO2 adapted cells of Chlamydomonas reinhardtii
DNA Res. 7 (5), 305-307 (2000)
20539644
11089912
                                                       Gaps
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/clone lib="Oryza minuta HybriZAP-2.1 XR library"
/note="organ: immature leaf"
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Unpublished (2003)
Contact: Jeong Sheop Shin
Plant Molecular Genetics
Graduate School of Biotechnology, University of Korea
1.16-701 Anam-dong 5/1 Seoul, Korea
Tel: 00 82 2 2320 928
Fax: 00 82 2 927 9028
     Length 544;
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ch 80.9%; Score 17.8; DB 1; 1
| Similarity 66.7%; Pred. No. 4.8e+02;
14; Conservative 5; Mismatches 2;
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66.7%; Pred. No. 4.8e+02;
iive 5; Mismatches 2;
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/mol_type="mRNA"
/db_xref="taxon:63629"
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Location/Qualifiers
                                                                                                    2 UGAACUCACUCGUGAGCUCTT 22
: |||:|| : ||:||||| || 197
197 TAAACTCATTCGTGAGCTCTT 477
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/mol_type="mRNA"
/mol_type="mRNA"
/mol_type="mRNA"
/mol="texton:3055"
/db_xref="texton:3055"
/db_xref="texton:3055"
/clone lib="C. reinhardtii CC-1690, Deflagellation
/clone lib="c. reinhardtii CC-1690, Deflagellation
/note="Vector: pBluescript II SK-; Site_I: ECORI; Site_2:
/note="Vector: pBluescript II SK-; Site_I: ECORI; Site_2:
Xho1; Deflagellation library, constructed by John Davies
and Jeffrey McDermott, combines cDNAs from CC-1690 cells
which had been re-synthesizing flagella for 15, 30 and 60
min after being deflagellated by pH shock. PolyA mRNA was
purified from each sample, pooled and cDNA synthesized.
The CDNA was directioned into lambda Zap II
(Stratagene) in the ECORI (5') and XhoRI (3') sites.
pBluescript II SK- plasmids were excised from the lambda
ZAP clones by superinfection with ExAssist (Stratagene)
phage. The library was normalized using method 4 described
in Bonaldo et al., (1996) Genome Research 6: 791-806."
/clone_lib="RPCI-23"
/note="Organ: Kidney/Brain; Vector: pBACe3.6; Site_1:
EcoRI, Site_2: ECORI; Female C57BL/6J mouse kidney and/or
brain genomic DNA was isolated and partially digested
with a combination of EcoRI and EcoRI Methylase. Size
selected DNA was cloned into the pBACe3.6 vector at the
EcoRI sites. The ligation products were transformed into
DH10B electrocompetent cells (BRL Life Technologies). "
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1030121E11.yl C. reinhardtii CC-1690, Deflagellation (normalized),
Lambda Zap II Chlamydomonas reinhardtii cDNA, mRNA sequence.
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Chlamydomonas raihhardtii
Bukaryota; Viridiplantea; Chlorophyta; Chlorophyceae; Volvocales;
Chlamydomonadaceae; Chlamydomonas.
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Grossman, A., Chang, C.-W., Davies, J., Harris, E., Hauser, C.,
Grossman, A., Chang, C.-W., Davies, J., Shifflow, C. and Stern, D.
Analyses of the Chlamydomonas reinhardtii Genome: A Model,
Unicellular System for Analyzing Gene Function and Regulation in
Vascular Plants. Project: 1030
Compublished (2002)
Contact: Charles Hauser
DCMB Box 91000
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                                                                                                                                                                                                                                                                     80.9%; Score 17.8; DB 8;
ilarity 71.4%; Pred. No. 4.8e+02;
Conservative 4; Mismatches 2;
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Fax: 919 613 8177
Email: chauser@duke.edu.
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Durham, NC 27708-1000
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Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.
E 1 (bases 1 to 547)
S Zhao, S., Nierman, W.; Feldblyum, T.; Malek, J.; Shateman, S.;
Akinret, B.; Levins, M.; Megann, S.; Tsegaye, G.; Geer, K.; Krol, M.; de Jong, P. and Fraser, C.M.
Mouse BAC End Sequences from Library RPCI-23
L Unpublished (1999)
Other GSss RPCI-23-114N6.TJ
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr.; Rockville, MD 20850, USA
Tel: 301 838 0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           547 bp DNA linear GSS 26-JUL-2000 RPCI-23-114N6.TV RPCI-23 Mus musculus genomic clone RPCI-23-114N6, genomic survey sequence.
                The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2: Arboi; The cDNA library was constructed from cells cultured in a carbon stream as acclimatized condition in which carbon dioxide concentration in the bubbling gas was changed from dioxide concentration in the bubbling gas was changed from
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Clones are derived from the mouse BAC library RPCI-23. For BAC
Clones are derived from the mouse BAC library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/orderingframe.htm)
or from Resea ch Genetics (info@resgen.com). BAC end page:
http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Seq primer: T7
Class: BAC ends.
                                                                                                                                                                                                                                                                                                      /clone="LC005h05_r"
/clone_lib="Chlamydomonas reinhardtii 5% to 0.04% CO2"
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                                                                                                                                                                                            /organism="Chlamydomonas reinhardtii"
/mol_type="mRNA"
/strain="C9"
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'organism="Mus musculus"
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strain="C57BL/6J"
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/clone="RPCI-23-114N6"
/sex="Female"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                db xref="taxon:3055"
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/note="Organ: Kidney/Brain; Vector: pBAC63.6; Site 1:
EcoR1; Site 2: EcoR1; Female C57BL/6J mouse kidney and/or
brain genomic DNA was isolated and partially digested
with a combination of EcoR1 and EcoR1 Methylase. Size
selected DNA was cloned into the pBAC63.6 vector at the
EcoR1 sites. The ligation products were transformed into
DH10B electrocompetent cells (BRL Life Technologies). "
                                                                                                              AZ695771 658 bp DNA linear GSS 24-JAN-2001
RPCI-23-240D2.TV RPCI-23 Mus musculus genomic clone RPCI-23-240D2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: szbaoœtigr.org
Clones are derived from the mouse BAC library RPCI-23. For BAC
Clones are derived from the mouse BAC library availability, please contact Pieter de Jong
(pdejong@mail.cho.org). Clones may be purchased from BACPAC
Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC end
pages: http://ww .tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Plate: 240 row: D column: 2
Seg primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               667 bp mRNA linear EST 01-AUG-200; Lambda Zap II.yl C. reinhardtii CC-1690, Deflagellation (normalized), Lambda Zap II Chlamydomonas reinhardtii cDNA, mRNA sequence. BQ820182 BQ820182. GI:22070844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear EST 01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                    Zhao,S., Nierman,W., Feldblyum,T., Malek,J., Shatsman,S.,
Akinret,B., Levins,M., Mcgann,S., Tsegaye,G., Geer,K., Krol,M., de
Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-23
Unpublished (1999)
Other_GSSs: RPCI-23-240D2.TJ
                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 658)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Shaying Zhao
Department of Bukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, USA
Tel: 301 838 0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:10090"
/clone="RPCI-23-240D2"
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482 TAAACTCATTCGTGAGCTCTT 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            228 GAGAACTCACTGGTGAGCTCT 208
                                                                                                                                                                                                                                             Mus musculus (house mouse)
Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="Female"
/lab_host="DH10B"
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                                                                                                                                                        genomic survey sequence.
AZ695771
                                                                                                                                                                                                 AZ695771.1 GI:12410231
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                                                              RESULT 14
AZ695771/c
                                                                                                                                DEFINITION
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BQ820182/c
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JOURNAL
COMMENT
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KEYWORDS
SOURCE
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/Mol type="mrkNA"
// strain="CC-1690 wild type mt+ 21gr"
// db_xref="texon:305s"
// db_xref="texon:305
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Chlamydomonas reinhardtii
Eukaryota, Viridiplantae, Chlorophyta, Chlorophyceae, Volvocales,
Chlamydomonadaceae, Chlamydomonas.
                                                                                                                                                        1 (bases 1 to 667)
Grossman, A., Chang, C.-W., Davies, J., Harris, E., Hauser, C.,
Grossman, A., Chang, C.-W., Davies, J., Harris, E., Hauser, C.,
Grossman, P., McDermott, J.D., Shrager, J., Slifflow, C. and Stern, D.
Analyses of the Chlamydomonas reinhardtii Genome: A Model,
Unicellular System for Analyzing Gene Function and Regulation in
Vascular Plants. Project: 1030
Contact: Charles Hauser
DCMB Box 91000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Pred. No. 5e+02;
5; Mismatches. 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Chlamydomonas reinhardtii"
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Job time: 1564.43 secs
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483 TAAACTCATTCGTGAGCTCTT 463
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: chauser@duke.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 UGAACUCACUCGUGAGCUCTT
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Tel: 919 613 8159
Fax: 919 613 8177
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Best Local Similarity 66.7%
Matches 14; Conservative
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